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GenCore version 5.1.4.p5_4578 copyright (c) 1993 + 2003 comp.gen_bbd.
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Run on:

April 29, 2003, 08.52.00°, Scarch time 14 Seconds (without alignments) 40.065 Million cell updates/sec

US-(9-647-749A-2

1 IKEYFTS 7 Perfect Score: Sednence:

Japop 60.0 , Gapext 60.0 OFFIG Serring table:

301532 segs, 80124803 residues Searched:

Word size :

Total combor of bits satisfying chosen parameter.

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

Database :

Published_Applications_AA:*

| Avan_E_G_proda_a3/2/pubgaa, TSSO6 (EW_PUB.pep:*
| Avan_E_G_P_PUBGAA, TSSO6 (EW_PUBG.pep:*
| Ava

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total sole distribution.

SUMMARIES

Result, SCC	Score 44	Query				
 	4.4	Matci	Match Length DB	DB	ID	Description
ा रच	•	57.1	6	1.6	US-09-277-074-44	Sequence 34, Appl
* - 1	,	57.1	10	ത	11 전 전 1 전 1 전 1 전 1 전 1 전 1 전 1 전 1 전	c)
7	*1	4.2.3	iO	ىد،	US-10-020-435A-5	sequence 5, Appli
	m	42.9	ī.	6	US-10-020-436A-14	14
u^	۲.,	42.3	u^	δ	US-09-911-129B-1+	13,
٩	3	42.3	ur.	1.0	US-09-859-214-42	Sequence 42, Appl
7	ب.	42.)	un	1.2	US-10-006-630-2	Sequence 2, Appli
σc	m	42.3	S	σ	US-09-727-963A-25	
Ġ.	æ.	42.3	٥.	10	18-720-868-60-80	Sequence 31, Appl
10	m	42.3	9	10	9-486-928-60-511	. Y
11	œ	42.3	Q	10	US-09-839-447A 'J	Ç
12	, 3 0	42.3	Ō	10	18-222-986-60-80	Sequence 31, Appl
13	٣	42.3	7	5	US-10-043-487-439	39,
14	т	4.2.	۲.	10	US-09-056-160B-5	~
1.5	cr.	42.3	7	10	US-09-056-160B-1.4	
16	m	42.3	r.	0	1 336.320.63.32	Sequence 7, Appli
17	m	42.3	7	ĵ	US-09-839-447A-51	Sequence 51, Appl
18	8	4.5	٢	<u>:</u>	1000-100-110-110-110-110-110-110-110-11	ä
51	m	42.9	7	10	US-09-898-461:17	Sequence 17, Appl

Sequence 19, Appl Sequence 19, Appl Sequence 20, Appl Sequence 8, Appl	25.	1.3	ज ज ज के लाग	Sequence set Applications of the Sequence 11. Application Applicat	Sequence 276, App. Sequence 19, App. Sequence 17, App. Sequence 16, App. Sequence 17, App. Sequence 17
	8 10 US-09-839-447A-50 9 12 US-10-66-151-16 9 12 US-10-152-654-2 9 12 US-10-172-654-3		0000 Hedd	10 US-09-872-832-38 10 US-09-305-928-25 12 US-10-066-151-21 8 US-08-945-038-11	10 9 (08-09-284-516A-276 10 9 (S-09-381-630-19 10 9 (S-09-791-389-16 10 9 (S-09-791-393-16 10 9 (S-09-791-393-16
20 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	48.85 48.85 48.85 48.85 50.00	90000 0000 0000 00000 00000	(ଅଟନ୍ତ (((((((((((((((((((40 41 43 44 45 46 33 462 465 465 465 465 465 465 465 465 465 465

ALIGNMENTS

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APELICAL INTERFECTION LINDS A.

FILLE E INVENTION IN THY DESIRATION OF TOMOR-SPECIFIC CYLLIXIO T DELLA
FILLE PERENENCE: SCREENS
TIERE PERENENCE: SCREENS
THERE OF THE CALL INTERPRETATION OF A STATE O
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APPLICANT: Sidney, John
TITLE OF INTERTION: 815 Mindles Poptidos and Their Uses
TILE REFERENCE: 996 82001321
CHEEN APPLICATION THREE TS C. 152,843A
CURRENT FILING DATE: 1995-05-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESOLT 2
US-08-472-843A-20
; Sequence 20, APELCALION US/ON452843A
...ont No. US20020098197A1
                                                                    7/99-277-074-34
Beginsen 34 Applitation PS/0927074
Padipath 6 RY (07197002820A)
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                                                                                                                                                                                                                                                                    GENERAL INFORMATIONS
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RESULT 1
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APPLICANT: CAMONT, ALAN
APPLICANT: WILLIAMS, DAVID
TITLE OF INVENTION: A METHOD FOR MAPPING THE ACTIVE SITES BOUND BY ENZYNES
TITLE OF INVENTION: A METHOD FOR MAPPING THE ACTIVE SITES BOUND BY ENZYNES
FILE REFERENCE: 179-54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: Description of Artificial Requence: Synthetic peptide US-10-020-436A-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 42.9%; Score 3; 1\Theta 9; Length 5; Best Local Similarity 100.0%; Pred, No. 2.7(-0.5); Matches 3; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 10,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            57.1%, Score 4, DB A
100 0%; Pred No 47*
tive 0; Mismatches
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FRICK APPLICATION NUMBER: US 08,7344, 814 PRICK FILINS DATE: 1994-11-23 PRICK APPLICATION NUMBER: US 08,7278, 8-44 PRICK ELLING DATE: 1994-07-11 NUMBER OF SEQ ID NOS: 30 SOFTHARE: FastSE2 for Windows Version 4.0 SEQ ID NO 20 LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              // Sequence 14, Application Us/10020436A
// Patent No. US20020155503A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: p53, 321-330
                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 100 (
Matches 4: Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-452-843A-20
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US-10-020-436A-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 EYF 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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. CIMER INFORMATION. Description of Artificial Sequence. Synthetic peptide US-10-020-436A-14
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TITLE OF INVENTION: INHIBITORS OF MADCAM-1-MEDIATED
INTERACTIONS AND METHODS OF USE THEREF H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 89
CORRESPONDENCE ADDRESS:
ADDRESSEE: Hamilton, Brook, Smith & Reynolds: P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
42.9%; Score 3: DB 9: Length F:
Bost Local Similarity 100.0%; Prod. No. 2.7e-05;
Matches 3: Conservative 0: Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                 Ouery Match 42.9%; Score 3; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 2.7e+05; Matches 3; Conservative 0; Mismatches 0; Indel:
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Sequence 13. Application BS/09411129B

Publication No. US20030040600A1

GENERAL HONOR US2003004060A1

APPLICANT Cleveland State University
APPLICANT Mano, Kenneth G.

TITLE 12 INVESTION. This mile 5. Contation inhibitor
FILE REFERENCE: 27433/04004

UNERHI FILING DATE: 2002-03 19

SOURMER HELD CALL CHOS: 17

SOFTWARE: Patentin version 3.1
APPLICANT: Schwender, Charles F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  // Sequence 42, Application US/09859214
// Futent No. US20020103111A1
// GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: Massachusetts
                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
ZIP: 02421
JOHEUTER REALABLE PARM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-911-129B-13
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1 EYF 4
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LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRI
                                                                                                                                                                                                             LENGTH: 5
                                                                                                                                                                                                                              TYPE: PRT
                                                                                                                                                                                                                                                                       FEATURE:
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Tue Apr 29 09:04:57 2003

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APPLICANT: Words and a statish APPLICANT: Requier, Fred APPLICANT: Requier, Fred APPLICANT: Regard, David APPLICANT: Williams, Revin APPLICANT: Williams, Revin APPLICANT: Afeyon, N. US20020150305Alta; APPLICANT: Paliwal, Sandeep APPLICANT: Paliwal, Sandeep TITLE OF INVENTION: Multi-dimensional molecular selection and analysis FILE REFERENCE SEPPIND CURFER APPLICANT SEPERATE SEPPIND CURFER APPLICATION NUMBER 32 10, 000, 500.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SCFTWARE: PatentIn Release #1.0, Varsion #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: /iabel- moditio-1 aa /note- "Leucine - NH2"
SEQUENCE DESCRIPTION: SEQ 1D NO: 42:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1
OTHER INPOPMATION: [label=mod)[fieltaa
/note="Ac - Leucine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 42.9%; Score 3; DB 10 L
Best Local Similarity 100 0%, Pred No. 2 75005;
Matches 3; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                              NAME: Brook, David E.
REGISTRATION NUMBER: 22,592
RERERENCE/DOCKET NUMBER: LKS95-12A.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781) 861-6240
TELEPRAX: (781) 861-6240
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS.
                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/109,879
FILING DATE: *UPARGNED
APPLICATION NUMBER: US 08/582,740
FILING DATE: 04 -JAN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                            APPLICATION NUMBER: US/09/859,214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2001-12-05
PRIOR PAPLICATION WINBER: 09/26/5, 993
PRIOR FILING DATE: 1999-03-12
PRIOR APPLICATION WIMMER 60/0000,518
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE: 1995-06-26
PRIOR AFELICATION NIMBER: 08/570, 570
PRIOR FILING DATE: 1996-06-26
PRIOR FILING DATE: 1996-08-86
                                                                                                                                                          FILING DATE: 16-May-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/EEY: Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Modified-site
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/10006630 Patent No. US20020150926A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 5 amino acids
                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
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2 FTS 4
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US-10-006-630-2
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Sequence 31, Application 08/0985072

Sequence 31, Application 08/0985072

Parent No. US20020025550A1

GENERAL INFORMATION.

SITIE OF INVENTOR. MILITAM M

STILE OF INVENTOR. MILITAM M

STREED OF INVENTOR. MILITAM M

STREED OF INVENTOR. MILITAM M

CORRENT FILING DATE: 2001-07-02

PRIOR PRIATATION NUMBER. 40,153,831

PRIOR FILING DATE: 1999-09-14

FRICE ALCIANTION DATE: 2000-09-14

FRICE ALCIANTION DATE: 2000-09-14

FRICE PRIOR FILING DATE: 2000-09-14

FRICE PRIOR DATE: 2000-09-14

FRICE PRIOR PRIOR DATE: 2000-09-14

FRICE PRIOR PRIOR DATE: 2000-09-14

FRICE PRIOR PRIOR SEC. 2000-09-14

FRICE PRIOR PRIOR DATE: 2000-09-14

FRICE PRIOR DATE: 2000-09-14

FRICE PRIOR PRIOR DATE: 2000-09-14

FRIC PRIOR DATE: 2000-09-14

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APPLICANT: V. I. Technologies, Inc.

TITLE OF INVENTION: METHED 'F IDENTIFYING A LIGAND FOR A TARGET MILETILE FILE REFERENCE: 18242-505

FILE REFERENCE: 18242-505

FRIGE RIT APPLICATION NUMBER: 18202/227,963A

CRREENT FILING DATE: 1999-12-02

NUMBER OF SECULATION HUBER: 1999-12-02

NUMBER OF SECULATION NON: 2.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         s ie:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90000
                                                                                                                                                                                                                                          FEATURE: OTHER INFORMATION: Description of Artificial Sequence, synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ) OTHER INFORMATION: Description of Artificial Sequence : peptibe corner information: ligand ms 09-727-969A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         o; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 42.9%; Score 3; DB 12; Length 5:
Best Local Similarity 100.0%; Pred. No. 2.7e+05;
Matches 3; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 42.4%; Score 3; DB 9; Length 5; Best Local Similarity 100.5%; Pred, No. 2.7e+05; Matches 3; Conservative 0; Mismatches 0; Indel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . Sequence 25, Application US/09727963A ; Patent No. US20020155106A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                           ORGANISM: Artificial Sequence
SOFIWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Bos taurus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-727-963A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-895-072-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-895-072-31
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                                         SEQ ID NO 2
LENGIH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGIH: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGIH: 6
                                                                                                                                                  TYPE: PRT
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                                                                                             LENGIH:
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Query Match 42.9%; Score 3; DB 10; Length 6: Best Local Similarity 100.0%; Pred. No. 2.7e+05; Matches 3; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0, Mismatches
; OTHER INFORMATION: Artificial Peptide US-09-839-447A-52
                                                                                                                                                                                            ; TYPE: PRT
; OFGANISM: Shigella Flexneri
US-10-043-487-439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3. Conservative
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Rest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                               US-09-986-552-31
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                                                                                              5 FTS 7
                                                                                                                          4 FTS 5
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LENGTH: 7
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                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 6
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                                                                                                                                                                   RESULT 12
                                          Ö
                                          Saps
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GENERAL INFORMATION:
APPLICATE SAILNEY MAITI
TITLE OF INVENTION: MEPATITE BY USUS CORE AND E ANTIGENS
TITLE OF INVENTION: HEPATITE BY USUS CORE AND E ANTIGENS
FILE REPERENCE: TRIPEP. 0200PL 0200PL
CURRENT APLICATION NUMBER: 05.09.7839.447A
CURRENT FILING FATE: 2001-06-09
PRIOR FILING DATE: 2000-04-21
NUMBER OF SEQ ID NOS: 111
SOFTWARE: FASISEQ for Windows Version 4.0
SEQ ID NO 52
LENTH: 6
                                                                                                                                             O. Indels
              Length 6:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 6;
             Query Match
42.9%; Score 3; LB Fr. Le
Best Local Similarity 100.0%; Pred. No. 2.7+-05;
Matches 3; Conservative 0; Mismatches 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42.9%; Score 3; EB 10: L
100 0% Pred to 2 7: 05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p Sequence 52, Application US, 73935447A pretent No. US20020058247A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Bast Foral Similarity 100 o
Matches 3, Conservative
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                                                                                   + FTS 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FEATURE:
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APPLICANT: CANFIELD, WILLiam
APPLICATION NUMBER: 05/09/086,552
CURRENT PRINTS DATE: 2001-11-09
PRIOR APPLICATION NUMBER: 05/05/36,872
PRIOR APPLICATION NUMBER: 05/05/36,831
PRIOR APPLICATION NUMBER: 05/09-08-14
NUMBER: 0F-09-14
NUMBER: 0F-09-14
SEQIENCE: 1999-09-14
SEQIENCE: NATION NUMBER: 05/09-09-14
SEQIENCE: NATION NUMBER: 05/09-09-14
SEQIENCE: NATION NUMBER: 05/09-09-14
SEQIENCE: NATION NUMBER: 05/09-09-14
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Sequence 479

Sequence 479

Sequence 470

Sequence 570

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42.9%; Score 3; DB 10; Length 7; 100.0%; Pred No 2.76.05; artive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                           COMPUTER PEROABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
COMPUTER: IBM PC compatible.
COMPUTER: SYSTEM: PC-DCS/MS-CDS
SOFTWARE: WINPALIN (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Baca, Manuel
APPLICANT: Wells, James A.
APPLICANT: Presta, Leonard G.
APPLICANT: Lowman, Henry B.
APPLICANT: Comman, Henry B.
TITLE OF INVENTION: ANTI-VEGF ANTIBODIES
NUMBER OF SEQUENCES: 131
                                                                                                APPLICANT: Baca, Manuel
APPLICANT: Wells, James A.
APPLICANT: Presta. Leonard G.
APPLICANT: Lowman, Henry B.
APPLICANT: Chen. Yvonne M.
FITLE GE TAVENTICH: ANTIBODILE
NUMBER OF SEQUENCES: 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER (35,59,0%,160B
FILTED DATE (06.Apr.1958
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60,054,856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PEGISTRATION NUMBER 28,616
REFRENCE/COCKET NUMBER: P1093R2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 124, Application US/09056160B Patent No. US20020032315A1
                                     // Sequence 5, Application US, 038681848
// Patent No. US20020042415A)
                                                                                                                                                                                                                                       COPPESPONDENCE AUGRESS
ADDRESSEE: Geneatech, Inc.
STREET: 1 DNA Way
CITY: South can Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genentech, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      South San Francisco
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 5:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 06-AUG-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
TYPE: Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Hasak, Janet E. PEGISTPATION NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Ue...
1 DNA Way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 3; Conserv
                                                                               GENFRAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                             94080
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US-09-056-160B-5
RESULT 14
HS-09-056-160B-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 FTS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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COMPUTER READALE FORM:
MEDIUM TYPE: 3 5 10-01, 144 Mb floppy disk
MEDIUM TYPE: 3 5 10-01, 144 Mb floppy disk
MEDIUM TYPE: 3 5 10-01, 144 Mb floppy disk
MEDIUM TYPE: 4 5 5 10-01, 144 Mb floppy disk
MEDIUM TYPE: MINDATA: (Genentech)
CUBRENT APPLICATION DATA: 424
MATCHAILANION MAREN: 424
MATCHAILANION MAREN: 6-0.051,856
MATCHAILANION MAREN: 6-0.051,856
MATCHAILANION MAREN: 8-0.6-051,856
MATCHAILANION MAREN: 8-0.6-051,856
MATCHAILANION MAREN: 8-0.6-051,856
MATCHAILANION MAREN: 124:
MARE: HEADAL TOWNING TOWNING THE MARCHEN FLOOR MAREN FOR SEQUENCE CANADAL MAREN: 650,952,988
MATCHAILANION FOR SEQUENCE CANADAL MARCHEN FOR SEQUENCE MARCHEN FOR SEQUENCE MARCHEN FOR SEQUENCE FOR SEQU
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GenCore version 5.11.4_p5_4579 Copyright (c) 1993 - 2003 Computed Ltd
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Run on

April 29, 2003, 04:45:15; Search time 20.833 Seconds (Without alignments) 23:673 Million cell updates/sec

US-09-647-749 1 LESYT 5 Title: Perfect score: : eduentes

Scoring table: OLIGO

Gapon 60 0 , Gapext 60 0

283224 seys, 96134422 residues Searched:

: ezise :

. .al number of hits satisfying chosen parameters

Minimum DB seq length: 0 Maximum DB seq length: 10

Listing first 45 summaries

pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		æ				
Result.	:	Query			;	
NO.	Score	Match	Match Length	98	QI	Description
-	c	60.0	6	7	PT0315	Iq heavy chain CRD
2	3	60.0	5	~	PT0324	. >
~	C1	0.0	-,	cı	A37832	phenol l-monocxyde
***	7		5	7	I39964	Вã
u"	(4	0.0	L)	C1	139966	
ý	(4	40.0	u)	c a	139965	libosomal protein
7	(4	40.0	L)	(4	I40469	BhaZX-like protein
œ	(4	40.0	L [†] ገ	CA	PT0308	iq heavy chain ORD
ć	(1	0.04	b"	۲.	01010	chain
10	2	40 0	ľ	~	-	T-cell receptor be
11	(4	40.0	'n	~	JT0870	phytosulfokine alp
2	CI	40.0	a,	`4	F11779	phosphoglycerate t
13	CI	40.0	e	C1	C 4	dihydrofolate redu
14	C4	40.0	9	্ৰ	B31263	dihydrofolate redu
15	7	40.0	¥	C1	TH0784	neuropeptide TR.6
16	2	40 0	7	, ,	XFYDGD	qalactose oxidase
17	64	○·0 *	t~	-	NTPG7	hypothalamic hepta
18	C1	40.0	۲.	C4	A60139	fatty-acid synthas
19	্ৰ	40.0	۲.	્ય	d'i	
୍ଟ	CI		r	C4	a i	wicilin 57F chair
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C4 C4	C4	40.0	r-	₹†	69	hypothetical L2 pr
53	C1	40.0	σ	C ŧ	0.1	glucose-6-phosphat
7.7	CI	40.0	œ	C4	B27867	homeotic protein U
25	2	40.0	œ	7	7	thymic humoral fac
42	7	0.04	10	्य	S.U.	eytochrome P450 AL
27	7	40.0	80	7	0.2	MHC class I histoc
28	2	0 0	œ	7	A25836	L-serine ammonia-l
ିଅ	C1	40.0	6	C4		phyllocaerulein -

Quory Match 60.0%; Score 3: DB 2: Length 9: Pest Local Shallarity 100.0%; Pred No. 2.86-05; Matches 7: Classiffe 0: Mismatches 0: Indels

2 ESY 4 |11| 6 ESY 8

5 QQ RESULT 3 A37832

fibrinoden beta ch Ig heavy chain CDB Ig heavy chain CRD Ig heavy chain CRD alphar2-marrowichou colligen alpha 2(V yrefraint 4 - bo fructuse-bisphasph 1 oxas 11 c34-rran Ig heavy chain V r I-cell receptor ga I-cell receptor ga I-cell receptor ga Gonadoliberin - sh conadoliberin - sh	FESULT 1 PTO315 FORTH 1 PTO315 FORTH 2 FORTH 2 FORTH 3	2; Length 9; 8c-05; 0; Indels · Japs · ·	PESULT 2 PT0324 If heavy thris, TRI? Legic (1917) - Commun (frighter) C.Specifics Holomosaplate (Late, 2018) C.Eate, 30.Septifies Berjacoperusylsten 30-Septifies Heavillate, 1918 C.Eate, 30.Septifies Berjacoperusylsten 30-Septifies Heavillate, 1918 C.Eate, 30.Septifies Berjacoperus, 1918 C.Eate, 30.Septifies Berjacoperus, 1918 C.Eate, 30.Septifies Berjacoperus, 1918 C.Eate, 30.Septifies Berjacoperus, 1918 C.Eate, 30.Septifies British Berjacoperus, 1918 C.Eate, 30.Septifies British
C24180 PT0231 PT0234 A6023 A60427 A60427 A60427 A1444 PF0443 G41946 G41946 G41946 A150463 KHSHG ALCAMENTS	dsion 30-Sep-1 dsion 30-Sep-1 dard, E.A.: Sh n of specific in d-109:47, Patt	Score 3: 18-2: Fred. No. 2:8e 0: Mismatchos	distoration of Sept. 1 distoration of Section of Sectio
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	inem promisers and services are services and services and services and services and services are services and services and services and services and services are services and services and services and services and services and services are	rit	1 3 % e 7 E 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
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онскасть встана фефф Онскасть встопия фы	RESULT 1 19 heavy c C; Species: 30 C; Date: 30 C; Cate: 30 C; Tates: 10 C; Tate: 10 C; Tat	Overy Match (0.0%) Best Local Similarity 100.0% Matches 3: Conservative Qy 1 LES 3 [11] Db 5 LES 7	RESULT 2 PT0324 T) h-11; C. Species C. Cator C. Cator C. Accessing F. Pamada F. Pamada F. Pamada A. Allice A. Molecule A. Molecule A. Reserved A. Molecule A. Reserved A. Molecule A. Reserved A. Reserved A. Molecule A. Reserved A. Reserved A. Molecule A. Reserved A. Reserved A. Molecule A. Reserved A. Reserved A. Reserved A. Reserved A. Molecule A. Reserved A. Rese

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Ryyamada, M., Wasserman, R., Reichard, B.A.; Shane, S.: Caton, A.J.; Kovera, G.
J. Exp. Med. 173, 395-407, 1991
A/Title: Preferential utilization of specific immunoqlobulin heavy chain diversity an
A/Reference number: PT0222: MJID:91108337: PMID:1899102
A/Accession: PT0308
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                                                                                                                                                                                                                                  ribosomal protein S4 - Bacillus megaterium (fragment)
C;Species: Bacillus megaterium
C;Species: Bacillus megaterium
C;Species: Bacillus megaterium
C;Accession: 13996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1946
C;Accession: 13996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1946
C;Accession: 13996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1946
A;Title: Characterization of the Bacillus subtills rpsD requlatory target site.
A;Recession: 139963, MCID-33015735; FMID:140026
A;Status: Preliminary: *rranslated from GR/FMRL/TDBJ
A;Roccession: 1399642; NID:444473
A;Cross-references: GB:M99042; NID:414473
C;Genetics:
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Cibate: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change lt-Aus-1995
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C:Species: Bacillus subtilis
C:Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 21-302-2003
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5, 2.8e+05;
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100.0%; Pred. No. 2.8
tive 0; Mismatches
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40.0%; Score 2; DB 2
Best Local Similarity 100.0%; Pred. No. 2.8
Matches 2; Conservative 0; Mismatches
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C:Keywords: heterotetramer; immunoglobulin
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Matches 2; Conservative
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    4 YT 5
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139965
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                                       C:Species: Pseudomonas Sp.
C:Date: 14-Jun-1991 #sequence_revision 14-Jun-199: #text_change 23-Jun-1993
C:Accession: A37832
C:Accession: A37832
B:PowLowski, J: Shingler, V.
J: Bacteriol. 172, 6834-6840, 1990
A:Title: In vitro analysis of polypeptide requirements of multicomponent phenol hydroxyl A:Accession: A37832; MUID:91072231; FMID:22:4259
A:Status: preliminary
A:Status: protein
A:Residues: protein
C:Seywords: oxidoreductase
phenol 2-monooxyqenase (EC 1.14.13.7) chain P5 - Freudomonas sp. (strain CF600) (fraymer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ridrindy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 6763-6770, 1992
A/Title: Characterization of the Bacillus subtills rpsD regulatory target site.
A/Reference number: 139963; MUE:93015735; PMID:1100226
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R.Grundy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 6763-6770, 1992
J. Sacteriol. 174, 6763-6770, 1992
A.Reference number: 139963; MUID: 93015735; PMID: 1400226
A.Accession: 139664
A.Status: preliminary; translated from GB/EMBL/CHAJ
A.Molecule type: DNA
A.Residues: 1-5 <RES>
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C.Date: 19-Jul-1996 #sequence_revision 19-Jul-1990 #text_change 19-Jul-1996
C.Accession: 139966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ribosomal protein S4 - Bacillus circulans (fragmen')
C:Species: Bacillus circulans
C:Date: 19-Jul-1996 #sequence_revision 19-Jul-1990 #text_change 19-Jul-1996
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0; Indels
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40.0%; Score 2; DB 2;
Best Local Similarity 100.0%, Pred. No. 2.8m-
Matches 2; Conservative 0; Mismatches
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A:Residues: 1-5 <RES>
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3.15

RESULT 4

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RESULT 5

139966

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Allydratelate relactors (EC 1.5.1.2), thymidylate synthase (EC 2.1.1.4), Hiasmodiu Cispecies: Plasmodium falciparum
Cispecies: Plasmodium falciparum
Cispecies: 12+169-1990 #sequence_revision 18+60-1990 #text_cnange 20-Mar-1996
Cisterion: A31269
Richerson, D.S., Walliker, D., Wellems, T.E.
Proc. Natl. Acad. Sci. U.S.A. 95, 914-9119, 1989
A/Tible: Evidence that a point mutation in dihydrofolate reductase-thymidylate syntha A/Reference number: A94217; MUID:89057886; PMID:2904149
A/Accession: A31263
A/Status: not compared with conceptual translation
A/Molecule type: DNA
A/Molecule type: DNA
A/Residues: 1-6 cePrix
C/Keywords: nethyltransferase; MADF: exidoreductase
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dlhydrothire policitor (pr. 1.5.) 3; / Plymilm; for exprinted (pr. 1.1.) 15; / Flismodiu
C.Speckes, Elasmodium fulliparum
C.Date: 28-Peb-1990 *sequence_revision 28-Peb-1990 *text_change 20-Mar-1996
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E,Feterson, D.S., Walliker, D., Wellems, T.E.
Proc. Natl. Acad. Sci. 0.s.A. 85, 9114-9118, 1988
A,Title: Evidence that a point mutation in dihydrofolate reductase togn.jylate syntha
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. 2.8e+05;
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100.0%; Pred. no.
''ve 0; Mismatches
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Jarity 100.0%; Pred. No. 2:8
Conservative 6: Mismatches
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Best Local Similarity 100 0%; Prad No.
Matches 2; Conservative 0; Mismatch
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Best Local Similarity
Matches 2: Conservat
                                                                                                                                  Best Local Similarity
A.Accession: JT0870
A.Molecule type: protein
A.Residues: 1-5 <MAT>
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2 LE 3
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                                                                                                         Query Match
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R:Matsubayashi, Y.; Sakaqami, Y.
P:Matsubayashi, Y.; Sakaqami, Y.
P:roc Nat! Acad Sct in A A 92 7622-7627 1900
P:Title: Phytosulfokine, sulfated peptides that induce the proliferation of single mesop A;Reference number: JT0870
                                                                                                                                                                                                                                                                                                           ig kappa chain V-III region (SD1) - human (fragment)
C)Species: Homo sap.ens (man)
C)Species: Homo sap.ens (man)
C)Acte: 2-Oct-1992 #Sequence_revision 23-Oct-1997 #text_change 16-Aug-1996
C)Accession: JT0520
R)Arker: R : Conley M F; Poillok R A
J EXP. Med. 169, 2.06-2119, 1989
A)Tible: Clonal diversity in the B cell repertoin of patients with X-linked agammadioby
A)Reference number. JT04[1, Mitp-A9274157; PMIC): 346547
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A.NOLE: the sequence shown here is one of eight productive V-D-J mu chain rearrangements
A.NOLE: a stop coden rerminates the sequence in 10 a V region
C.Keywords: heterotetramer; immunoglobalin
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A:Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A:Reference number: PT05.04: MUID-412776.01; PMTD-1-1185.8
A:Accession: PT061.0
This: translation not shown
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Cispecies: Mus musculus (house mouse)
Cispecies: Mus musculus (house mouse)
Cispate: 17-Jul-1992 #saquapea=_regisien 17-Jul-1992 #text_change 30-May-1997
B.Peeney, A 3 3 4 5 15-124, 1991
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                      40.0%; Score 2: DB 2: Length 5: 100.0%; Pred. No. 2.80.05; Live 0; Mismatches 0; Indel:
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A.Experimental source: newborn thymus, strain BALHVC
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C.Species: Asparagus officinalis (garden asparagus)
C.Date (D.Apri2002 #sequence_revision 02-Apri2000 #
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Matches 2; Couservative
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Sectes: Ascaris suum (pig roundworm)

Jecles: In-hn-1943 *sequence_revision 10-Jun-10-0 *text_change 28-Apr-1995

C.Accession: JH0784 *C.; Curry, W.J.; Johnston, C.F.; Thim, L.; Halton, D.W.; Buchanan, K. Blochem. Biophys. Ros. Commun. 187, 1333-1339, 1902

Alitle: The primary structure of TE-6: a novel : iropeptide from the nematode Ascaris & A. Recession: JH0784 *MDID-93038603; PMID: i117808

A. Accession: JH0784 *MDID-93038603; PMID: i117808

A. Molecule type: protein

A. Residues: 1-6 <SMA>

Residues: 1-6 <SMA>

A. Residues: a novel : iropeptide

C. Keywords: neuropeptide
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A;Reference number: A94217; MUID:89057886; PMID:2:4149
A;Accession: B31263
A;Status: not compared with conceptual translatic:
A;Moleus: hot compared with conceptual translatic:
A;Moleus: 1-6 <PET>
C;Keywords: methyltranslerase: NADP: oxidoreducta::0
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GenCore version 5.1.4 p5_4578 Copyright (\sigma) 1993 - 2003 Complien Ltd.
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Run on.

April 29, 2003, 08:30:44; Search time 11.6667 Seconds (without alignments)

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112892 seqs, 41476328 residues Searched:

size

Total number of hits satisfying chosen parameters

Minimum DB seq length, 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

Miller Charles and Sec. SW18SProt_40:*

Presentation is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total same distribution.

SUMMARIES

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Query Match 40.0%; State 2: DB 1: Length 5: Best Local Similarity 100.0%, Fred. No. 1.18+05; Matches 2: Classration, 7: Mishatches 2: Classration, 7: Mishatches 2: Local

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21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last Sequence update)
21-JUL-1986 (Rel. 01, Last annotation update)
Hypothalmic heptapeptide.
Sus schola (Pig)
Enkatycta, Metazca, Chondata, Cianiata, Vertebrata, Enteirostomic:
Mammalia, Entheria, Cetarliodactyla; Suina; Suidae; Sus.

7 AA

PRT;

STANDARD;

HY7_PIG F01153;

RESULT D HY7_PIG

P82072 Litoria nub P82073 Litoria nub P19472 escherichia P82028 cea mays (m P28055 chamaydia t P28055 tautubacill P1735 mytilus edu P1491 locusta mid P42485 leftinctars P44495 sarrothaga P44495 sarrothaga	ytusaifukine-beta (PSK-	Dates careta (Carrot). Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta: Bukaryota, Viridiplantae, Streptophyta, Embryophyta; Tracheophyta: Asteridae, wasterids II; Apiales, Apiaceae; Dancus, NCBLITAXID=4039; SEQUENCE, AND IDENTIFICATION BY MASS GFECTROMETRY. STRAIN-CH. US-Haidmakiqusun; STRAIN-CH. US-Haidmakiqusun; Manada H., Matsun T., Vamamoto M., Matsubayashi Y., Robayashi T., *A secreted peptide growth factor, phytosulfokine, acting as a stimulatory ractor of carrot somatic embryo formation."; Plant Cell Physiol 41-2-3(200). *I SAKGAMA OF CATOR SOME OF THE EMPROPERMINATION INTO THE SCMATIC EMBRYOS. **I FUNCTION IN PERSENTE OF 14-C, SIIMULAIES FALIFERALIS OF THE CELLS. But I GOES NOT SIIMULAIE DIFFERNIATION INTO THE SCMATIC EMBRYOS. **I SUBCELLULAR LOCATION: Secrete MY SIMILARITY). **I SUBCELLULAR EMBRANE REDEPTOR (BY SIMILARITY). **I SUBCELLULAR EMBRANE REDEPTOR (BY SIMILARITY). **I SULFATION IS SULFATION. **PROPTIDE MANAGEMENT OF THE PHYPOSULFORINE FAMILY. **Growth factor of SULFATION. **SULFATION. **SULFATI
RE32_LITRU RE32_LITRU TRN3_ECCLI UC22_BAIZE UXA4_CHLTR ACPH_RABIT ASP2_LACSN ACPL_LACNT OVM_LEPDE IMOF_SARBU TRNF_ESPEU	ALIGNMENTS ANDARD: PRT: 5 AA. 40, Greated) 40, tast sorpoone update) 40, Last annotation update) 64, Last annotation update)	Detail. Datus carota (Carrot). Datas and a displantae, Streptophyta, Embryophyta; Tra Spermatophyta; Magnollophyta; endlocyledons; core endic Asteridae; wasserids II; Apiales; Apiaceae; Dancus. NCBI_TaxID=4039; SECUENCE, AND IDENTIFICATION BY MASS SPECTROMETRY. STRAIN:
00000000000000000000000000000000000000	STANDARD; (Rel. 40, Grea (Pol. 40, Last (Rel. 40, Last Useralpha (FEE:	a (Carrot). Airidiplantae. a: Magnollophy 1039; 1019: Airidiplantae. 1029: Airidiplantae. 1020: Airidiplantae. Airidi
40000000000000000000000000000000000000	PESULT 1 PSK_DAUCA TID PSK_DAUCA AC PS8261; DT 16-OCT-2001 (Rel. 40, 10, 10, 10, 10, 10, 10, 10, 10, 10, 1	Dates); Dates carot Bukaryota, V Stermatophy Asteridae, IV STATING, AN STATING STA
	RESULT PSK_DAN TO DA AC DA DT DI DT DI DT DI	

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Eukaryota; Melazoa; Arthropoda, Mandibulata, Fancrustacea; Crustacea;
Malacostraca; Eumalacostraca; Eucarida; Decapoda; Pleocyemata;
Brachyura; Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBI_TaxID=6759;
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                                                                    MEDLINE-81213990 PubMed-b.20.578; Chang R.C. Coy D.H., Chang R.C. G. Huang W.-Y., Arimuta A., Kedding T.W., Coy D.H., Saffran M. Novg A. Hamilton J.W., Cohn D.Y., Schally A.V., affran M. Koyg A. Hamilton J.W., Cohn D.Y., Schally A.V., "Isolation, structure and synthesis of a hep-papehide with in vitro ACIH-releasing activity from porcine hypothalamus."; Pirm. Metab. Res. 13:228-232(1981). PIR; A01417; NYPG7.
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01-007-1994 (Rel. 30, Last annotation update)
01-o07-1994 (Rel. 30, Last annotation update)
Dactylium dendroides (Cladobotryum dendroides)
Bukaryota, Enngi, Assomycota, Pezisomycotin, Sordaricmycotes;
NGBLTaxID=5132;
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llarity 100.0%; Pred. No. 1.le+05;
Conservative 0; Mismatches 0; Indels
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MEDLINE-98121193; PubMed-9461295;
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                                                           SEQUENCE, AND SYNTHESIS.
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Matches 2; Conserv
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Electropheresis 13:707-714(1992).
-:- MISCELANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 4 9 ITS MW IS-65 kDa SWISS-2DPAGE; P30087; HUMAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Macaca fuscata fuscata (Japanese macaque).
Eukaryota, Metazoa; Chordata; Craniala; Vertebrata; Euteleostomi:
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi:
Mammalia; Eutheria; Primates; Catalihini, Hominidae, Homo.
NCFI_TaxID=4606;
      "Isolation and identification of multiple neuropeftides of the allatostatin superfamily in the shore crab Carcinus maenas."; Eur. J. Bicchem 259,727-734(1997).

-: FUNCTION: MAY ACT AS A NEUFOTPANSMITTER OF NEUROMODULATOR.

-: SIMILAPITY: PELONGS TO THE ALLATOSTATIN FAMILY.

Neuropeptide: Amidation; Multipene family.

MOD_RES 8

AMIDATION.
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15-JUN-2002 (Mel. 41, Last annotation update)
Fibrinogen beta chain (Contains: Fibrinopeptide B) (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hughes G.J., Frutiger S., Faquet N., Ravier F., Fasquali C.
Sanchez J.-C., James R., Tissot J.-D., Bjellqvist B.,
Hochstrasser D.F.;
                                                                                                                                                                                                                                                                                                                                                                            01-APR-1993 (Rel. 25, Created)
01-APR-1993 (Rel. 25, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Unknown protein from 20-page of plasma (Spot 2) (Fragment)
Homo sapiens (Human).
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                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Pred No. 1.1e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 AA; 944 MW; 0017720455BB06DA CRC64;
                                                                                                       MOD_RES 8 AMIDATION. SEQUENCE 6 AG 613 NW, TOD8064:FABIT6878 OFC64:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=93092937: PhrMcd=1459097;
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                                                                                                                                                                                                                               3 SY 4
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Thorpe A.;
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Nakamura S., Takenaka O., Takahashi K.;
"Fibrinopeptides A and B of Japanese monkey (Macaca fuscata) and patas monkey (Erythrocebus patas): their amino acid sequences, restricted mutations, and a molecular phylogory for macaques, quenons, and baboons.";
J. Blochem. 97:1487-1492(1985).
-1- FUNCTION: FIRRINGER HAS A DOUBLE FUNCTION: YIELDIN: MONOMERS THAT POLIXMERIZE, INTO FIRRINGHAS A CLIMACTOR IN PLATELET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              the git of West Indies Sugarizate Frutstalk Errett Wegvil (Diapreparabbreviatus L.) larvae. ".

Comp. Blochem. Physiol. 118E-861-867(1397).

-!-CATALYTIC ACTIVITY: Bandom hydrolysis St. (4-alpha-D-galactosiduronic linkages in pectate and other galacturonans.

-!-INDUCTION: INHIBITED BY CITRUS PGIP.

-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS PROTEIN IS:
                                                                                                                                                                                                                                                                                                                                                                                (ALPHA, BETA AND GAMMA), LINNED TO EACH SCHEE MY DISULFILE BONDS.

MISCELLANDSOLS CONVERSION OF FIRELWORD TO FIRELY IS TRIGGERED BY THROMEIN, WHICH CLEAVES FIRELNORPHIDES A AND B FROM ALPHA & BETA CHAINS, AND THIS EXPOSES THE N-TERMINAL FOLVMERIZATION SITES RESPONSIBLE FOR THE FORMATION OF THE SOFT SLOT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE-Larval dut; Doostdar H.; Mayer R.T.; boostdar H., McCollum T.G., Mayer R.T.; "Purification and characterization of an ends polygalacturonase from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Herry of the Ass.
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15-302.1998 (Rel. 36, Last sequence update)
15-301.1998 (Rel. 41, Last annotation update)
15-501.2002.002 (Rel. 41, Last annotation update)
15-501.2003.002 (Rel. 41, Last annotation update)
15-501.2003.002 (Rel. 41, Last annotation)
15-501.2003.002 (Re
                                                                                                                                                                                                                                                                                                                                                      -1 - SUBUNIT: HEXAMER CONTAINING 2 SETS OF A NONLUBENTICAL CHAINS
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-!- SIMILARITY: WEAK, TO OTHER POLYGALACTUR WASES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERPTO, IPRO0181; FIBRINGGELC.
PROSTTE: PSO0514; FIRBIN_AG_C_DOMAIN; PARTIAL
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Mismatches
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Non TER
MEDLINE-85289140; PubMed-3928610;
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                                                                                                                                                                                                                                                                                                                        AGGREGATION.
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P81179;
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01-301-1993 (Ref. 26, Last sequence update)
01-301-1993 (Ref. 26, Last sequence update)
01-301-1993 (Ref. 1993 Property of Last annotation update)
03-1994 (Ref. 1994 Property of Liver (Spot 113) (Fragment).
03-1995 (Ruman).
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MEDINE-9203297; PubMed-7895722;
Corbert M. Wheeler C.H., Raker C.S., Yacoub M.H., Dunn M.J.,
"The human myocardial two-dimensional gel protein database: update.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Electrophoresis 15:1459-1465(1994).
-!- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKN.WW
                                                                                                                                                                                                                                                                       Homo sapiens (Human).
Eukaryota, Metaroa: Chordata, Craniata, Vertebrata; Euteleostomi:
Mammalia: Eutheria: Primates: Catarrhini; Hominidae; Homo.
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Mammaila, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                      01-FBB-1995 (Rel. 31, Created)
01-FBB-1995 (Rel. 31, Last sequence update)
15-02T-2001 (Rel. 40, Last annotation update)
Unknown protein from IP-page of heart (Spot 5603) (Fragment).
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Rest Local Similarity 100.0%; Pred. No. 1.1
Matches 2; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROTEIN IS: 6.0, ITS MW IS: 55.3 kDa.
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MEDLINE=94147969; PubMed=8313870;
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Matches 2) Conservat
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                                    UHA2_HUMAN
P40929;
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UHA2_HUMAN
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"Two-dimensional gel electrophoresis separation and N-terminal "Two-dimensional gel electrophoresis separation masteurianum W5.";
sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:06.2966(1998).
.: MISCELLANEDUS. ON THE 2D-3EL THE PETERMINED F! OF THIS UNKNOWN PROTEIN IS: 6.5, ITS MW IS: 38.0 kDa.

VARIANT 8 8 M -> D.
NON TER 9 9
                                                                                                                                                                                                                                                                                                                                                                                       IISSUE-Liver;

MEDLINE-9447969; PubMed-8313870;

Hughes G.J., Fruitiger S., Paquet N., Pasqual, C., Sanchez J. C.,

Hughes G.J., Fruitiger S., Paquet N., Hochstrusser D.F.;

Tissot J.-D., Bairoch A., Appel R.D., Hochstrusser D.F.;

"Human liver protein map: update 1993.";

Electrophoresis 14:1216-1222(1993).

-:- MISCELLANROGS: ON THE 25-3EL THE LEIERMINIC FI OF THIS CHRICHN PROTEIN IS. 5.5, ITS MM IS: 12 KDa.

SWISS-2DPAGE; P31934; HUMAN.
                      01-JUL-1993 (Rel. 26, Created)
1-JUL-1993 (Rel. 26, Last Sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
Unknown protein from 2D-page of liver tissue (Spot 94) (Fragment).
                                                                                                                                                                      Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Verteurata, Euteleostomi;
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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100.0%; Pred, No. 1.10+05;
tive 0; Mismatches 0; Indols
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/FTId=VAR_0000.1.
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18-30L-1998 (Rel. 36, Last sequence update)
18-30L-1998 (Rel. 36, Last annotation update)
Unknown protein 3P 19 from 20-page (Fragment
Clostridium pasteurianum.
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Matches 2; Conserv
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P81355;
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SEQUENCE
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Wabnitz P.A., Bowie J.H., Tyler M.J.; "Caerulein-like peptides from the skin glands of the Australian Plae montains tree frog Litoria citropa, Part 1. Sequence determination
p81731, 30-MAY-2000 (Rel 34, Created) s0-MAY-2000 (Rel 34, Created) s0-MAY-2000 (Rel. 34, Last annotation update) s15-UN -2002 (Rel. 41, Last annotation update) Mainopeptidase N (EC 3 4 11 2) (CryIA(C) receptor) (Fragment). Helicoverpa armigera (Cotton bollworm). solutions armigera (Cotton bollworm). Sinsecta, Perryota, Arthropoda; Mandibulata, Fauctustacea, Bekapeda; Ditrysia; Noctuoidea; Noctudae; Heliothinae; Helicoverpa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SPECIES=L.splendida:
MEDLINE=20069471: PupMed=10601876;
Wabblitz P.A., Bowle J.H., Tyler M.J., Wallace J.C., Smith B.F.:
Wabblitz P.A., Bowle J.H., Tyler M.J., Wallace J.C., Smith B.F.:
"Differences in the skin peptides of the male and female Australia:
tree frog Litoria splendida. The discovery of the aquatic male sex
pheromone splendipherin, together with Phe8 caerulein and the
antibotic peptide caerin 1.10.";
antibotic peptide caerin 1.10.";
Eur. J. Biochem. 257.269-27(2000).
:- PUNCTION: HYPOTENSIVE NEUROPEPTIDE (PROBABLE).
                                                                                                                                                                                                                                                                                                                                                                                                              fnglc S.S., Trivedi N., Prasad R., Rao K.K., Chatpar H.S.;
"Aminopeptidase-N as a receptor for Bacillus thuringlensis CryIA.'
toxin from Helicoverpa armigera.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Litoria citropa (Australian blue mountains tree frog), and
Litoria splendida (Magnificent tree frog).
Enkaryota: Metaroa: Chordata: Craniata: Vertebrata: Enteleostori:
Anghilkia: Battarbia: Anora: Nechatrachia, Pafoncilea: Hylilae:
Pelodryadinae: Litoria: Anora: NCBI_TAXID=94770, 30345;
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o. 4.3e+03;
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Interpro: IFROGOLNG: Zn_MTpeptdse.
PROSITE: PSOG142: 71NC_PROTEASE: PARTIAL.
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16-00T-2001 (Rel. 40, Last annotation update)
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100.0%; Pred. No.
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es 2: Conservat
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MEDLINE-97374000; PubMed-9230483;
Steinborner S T , Waugh R J , Rowie J H Wallage J C , Tyler M J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Litoria xanthomera (Orange-thighed frog).

Bukaryota, Metazoa, Chordata, Graniata, Vertebrata, Euteleostomi;
Pmphibia: Batrachia, Anura, Neobatrachia, Hubencidea, Hylidae,
NCBL_TaxIb=79697;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "New Caerin antibacterial peptides from the skin glands of the Australian tree frog Litoria xanthomera.";
J. Pept. Sci. 3:181-185(1997).
I. FINATION HYPOTENSIVE NUTROPEPTIDE
I. TISSUE SPECIFICITY: SECRETED BY THE SKIN SORSAL GLANDS.
I. MASS SPECIFICITY: MW-1354; MRTHOD=FAB.
I. SIMILARITY: BELONGS TO THE GASTRIN/CHOLESSTOKININ FAMILY.
InterPro; IPR001651; GASTRIN; 1.
                                                                                        OULFAPER SPECIFICMETRY: MM-1366; METHOD-Electrospiay.
1- SIMILARITY: BELONGS TO THE GASTPHOGHALFON-DAKININ FAMILY INTERPROPER SPECIFICATION.
FORESTE: PROG1651; GASTPIN: FALSE_NEG.
Amphibian Skin: Hypotensive agent: Amidation. Sulfation.
MOD PES.
1 FYREGIE: PARTE ACID.
-i- TISSUE SPECIFICITY: SECRETED BY THE SKIN (**PSAL GLANDS.) -i- PTM: ISSPS/PM 1 2Y4 (*IFFESS FP/M ISSPS/PM ) ; IN NOT BEIN)
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MOD_RES 1 PYRROLIDONE CAFR XYLIC ACID.
MOD_RES 4 SULFATION.
MOD_RES 10 AMIDATION.
SEQUENCE 10 AA: 1290 MM; 99DBF3837861HR4A CRC64;
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        MOD_RES
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        AMIDATION.

        SEQUENCE
        10 AA.: 150C MW: 990BFCC27861EPF A. 18024;
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15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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DE (II) Hebrita II).

S Allivator mississippiensis (American alligator).

OC Bukanyota, Metazaa, Chordata, Craniata, Verrebrata, Euteleostomi, OC Bukanyota, Metazaa, Chordata, Craniata, Verrebrata, Euteleostomi, OC Bukanyota, Metazaa, Chordata, Craniata, Verrebrata, Euteleostomi, OC MITELIATID=R466.

RN HTLLATID=R466.

RN FELTINS-19152336, BubMed-1882082;

RN EGINNS-9152336, BubMed-1882082;

RN ELONE-9152336, BubMed-1882082;

RN ELONE-9152336, BubMed-1882082;

RN ELONE-9152336, BubMed-1882082;

RN ELONE-9152336, BubMed-1882082;

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April 29, 2003, 08.35.29 , Search time 19.5832 Seconds (without alignments) 52.60% Million cell updates/sec Run on:

1 LESYT 5 sequence: Scoring table: OLISO Gapext 60 0 Gapext 60 0

671580 seqs, 206047115 residues

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sp_rodent:*
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sp_vertebrate:* sp_mhc:* sp_organelle:* sp_phage:* sp_plant:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total sector distribution. sp_rvirus:* sp_bacteriap:* sp_archeap:*

sp_unclassified:*

SUMMARIES

Description	Q15898 homo sapien	P72221 pseudomenas	Q9r3x0 planktothri	Q9y4j4 homo sapien	ú9y4j3 homo sapien	Of0773 home sapted	Q94623 manduca sex		boophil	gadana iocusta mig	diadema	Q8wAq6 diadema mex	Q8w8g5 diadema ant	Q8w8g4 diadema mex	ū8w8g3 diadena pau	Q8w8g2 diadema sav
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ALIGNMENTS

015898	868			
QI	015898	PRELIMINARY;	PKT:	8 AA.
AC	015898;			
ΕŢ	01-NOV-1996	(TremBLrel. 01,	Created)	
DI	01-NOV-1996		Last seque	Last sequence update)
E	01-DEC-2001 (TrEMBLiel.	(Trembliel, 19.	Last annot	Last annotation update)
DE	(clone XP6Al	(clone XP6AllB) (Fragment).		
S	Homo sapiens (Human)	(Human).		
၁	Eukaryota, M	etazoa, Chordata	u Craniata	Bukaryota, Metazoa, Chordata, Graniata, Vertobrata, Buteleostomi:
ટ	Mammalia; Eu	theria; Primates	:: Catarrhi	Mammalia; Eutheria; Primates; Catarrhini; Hominidue; Homo.
×	NCBI_TaxID=9606;	606;		
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R.P.	SEQUENCE FROM N.A.	M N.A.		
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8	Lee CC., Y	azdani A., Wehne	ort M., Bai	
ΚĀ	Coolbaugh M.	I., Chinault C.A	A., Raldini	Coolbaugh M.I., Chinault C.A., Raldini A., Lindsay E.A., 25a - 3. T.
RA	Caskey C.T.H.;			
RT	"Isolation o	f chromosome-spe	scific gene	s by reciprocal probles of
E.T.	arrayed cDNA	s and cosmid lib	praries.";	arrayed cDNAs and cosmid libraries.";
R.	Hum. Mol. Ge	Hum. Mol. Genet. 0:0-0(1995).		
0: (1	EMBL; 132078	EMBL; 132078; AAA74888.1;		
F.	NON TER			
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Ç.	ы	8 AA; 528 MK; 3	84A415B0477	34A415B0477B45BB CRC64;
<u>ं</u>	Query Match		Score 3;	60.0%, Score 3; DB 4; Length 8;
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BUELINE-932-9200. FubMed-8330113.

MAKI H., Mori A., Fujiyama K., Kinoshita S., Yoshida T.;

"Ciouing, sequence analysis and expression in Escherichia coling and expression in Escherichia coling agene encoding an adjainate lyase from Pseudomonas sp. OS-ALG-9.",

Gen. Microbiol 13-987-993(1993)

EMBL: D38469; HAA21704-11;
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01-MAY-2000 (TYENNELO] 13 Last sequence uposte)
(1-MAY-2001 (TERNELO) 16 Last annotation spate)
Ribulose-1,5-bisphosphate carboxylase/oxygenuse large subunit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (will-1994) to the BMRLyNeshank/Ness databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bacteria, Cyanobacteria, Oscillatoriales, 71 offictbrix
01-FEB-1997 (TrEMBLrel, 02, Created)
01-FEB-1997 (TrEMBLrel, 02, Last sequence uplate)
01-PET-2001 (TrEMBLrel, 19, Last annotation indate)
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                                                                                                                                                                                       Pseudomonas sp. (strain us-ALG-9).
Bacteria, Proteobactelia.
NCBI_TaxID-86018;
                                                                                                                    Alginate lyase (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
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Best Local Similarity
Matches 2, Conserv.
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                                                                                                                                                                                                                                                                                                                                                                                  SIRAIN-OS-ALG-6;
                                                                                                                                                                                                                                                                                                                                                                                                             Fujiyama K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Fragment).
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RESULT 4

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TISSUP-BONE MAPPOW.
TISSUP-BONE MAPPOW.
TIGHE UP-ACTOR TO PERMODE TO 19324;
Tighe Up-ACTOR TO 19324;
Tighe Up-ACTOR TO 19324;
"Alternative (a) chromesome in the t(8:2) of acute mysicial leavemia.
                                                                                                                                                                                                                                                                                                               SECUENCE FROM N.A. MEDLINE-95002916; PubMed-7919324; Fighe J.E., Calabi F.; "Alternative, out-of-frame runt/MTG8 transcripts are encoded by the delivative, (6) chromosome in the 1(8.21) of 1.11e myelvil 12.1ection M2.".
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Enbary ta, Metapoa, Chen Mata, Chaniata, Vertetrata, Estelesstomi:
Mammalla: FutherMa: Primates: Cataribini, Hominidae; Homo.
                                                                                                                                                                                                          Enkaryota: Metazona: Chordata: Cruniata: Vertebrata; Euteleostomi:
Mammalia; Eutheria: Primates; Catarrhini; Hominidae: Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                40.0%: Score 2: DB 4: Length 8:
llarity 100.0%; Pred. No. 6.7e+05;
Consermative 0: Mismatches 0: Indels
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
                                                    ñi-NnV-1999 (TrEMBLrel. 12, Created)
01-NoV-1999 (TrEMBLrel. 12, Last sequence update)
01-NoV-1999 (TrEMBLrel. 12, Last annotation update)
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01-MAY-2000 (TrEMBLrel 13, Last sequence update)
                                                                                                                                         Ront/82nt/MTG8 protein (Fragment).
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01-NOV-1999 (TrEMBLrel. 12) Last an
hort-Sear ATGA protein (Fragment).
FintT-Sear ATGA protein (Fragment).
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EMBL: S74094; AAD14973.2; -.
NON_IER 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pi-nd P4 2:18-2:2:(1094).
EMBL: S74092: AAD14144.1; -.
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PRELIMINARY;
                                                                                                                                                                                                       Homo sapiens (Human).
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Best Local Similarity
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Matches 2, Conserma
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MEDLINE-97165493: PubMed=9013254;
MEDLINE-97165493: PubMed=9013254;
Jindra M. Huang J.Y. Malone F., Asahina M. Riddiford L.M.;
Jindra M. Huang J.Y. Malone F., Asahina M. Riddiford L.M.;
Indentification and mRNA developmental printing of two ultraspirations in the epidermis and wings of Manda a sexta.";
Insect Mol. Blol. 6:41-53(1997).
ENBL; US7921, AAF4222 1.
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Bebrayyota: Metaroa: Arthropoda; Tracheata: Hovapoda: Inserta;
Pterygota: Moptera; Endopherygota: Lepidoptora: Glossata; Ditrysla:
Sphinglodea: Sphinglidae, Sphinglinee, Manduca:
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Eukaryota, Metazoa, Chordata, Craniata, Vertibrata, Euteleostomi;
Mammalia; Butheria: Carnivora; Fissipedia; Unidae; Canis.
NCBL_TaxID+9615;
                                                              Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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O
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                                                                                                                                                 Ho S.;
Submitted (DEC-1999) to the EMBL/GenBank/DDFG databases
                                                                                                                                                                                                                                                                                                                                                                                                                                             01-FEB-1997 (TrEMBLrel, 02, Created)
01-FEB-1997 (TrEMBLrel, 02, Last sequence update)
01-NoV-1998 (TrEMBLrel, 08, Last annotation update)
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                   8 AA
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                  FF126G10.1 (Parvalbumin) (Fragment)
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                                                                                                                                                                           EMBL, 18184, CABO5099 3,
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                                               Homo sapiens (Human)
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Matches 2; Conserva
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                                                                                              NCBI_TaxID=9606;
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6 ES 7
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                                 PVALB.
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NEDITIE=7922731, F.P.We3.10368951;
Campbell N. U. Barker S.C.
The novel mitted-noutial year arrangement of the cattle tick.
F.-Phillus mitting of the F.13 tradem reportation of a calling region.";
Mol. Biol. Evol. 16:732-740(1999).
EMBL; AFI10616; AAD28386.1; -.
                           MEDEINE=97049323; Pubmed-8894053;
Venta P.J., Brouillette J.A., Yuzbasiyan-Gurkan V., Brewer G.J.:
"Gene-specific universal mammalian sequence-tagged sites: application
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sdet
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                                                                                                                                                                                                                                                              Venta P.J., Gao Y , Alexander L , Yuzbasiyan-Gurkan V.; "Dinucleotide repeat polymorphism in the canine retinoblastoma (RBI) gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryotta; Metazoa; Arthropoda; Chelioerata; Arachnida; Acari;
Parasitiformes; Ixodida; Ixodidae; Boophilus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota, Metazoa, Arthropoda, Trachesta, Hexapoda, Insecta,
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                                                                                                                                                                                                                                                                                                                                                           Submitted (JMN-1999) to the EMBL, MenPank, MoRA databases.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
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01-NPV-1966 (IDEMELRE). 01. Created)
01-NPV-1966 (IDEMELRE). 01. Last sequence update)
01-NPC-2001 (IDEMELRE). 19. Last sequence update)
01-PEC-2001 (IDEMELRE). 19. Last sequence update)
01-PEC-2001 (IDEMERRE).
Electron migrateria (Migratery locust).
Micochondrion.
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SPUGENCE NIAA ++0 100, SPHG441)4PETSKPG IPPE44,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              NON_TER 8 8 8 SPQUENCE 8 AA; 945 MW; 1425PR19676721E3 CRC64;
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                                                                                                                                                              Biochem. Genet. 34:321-341(1996).
                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AF155737; AAD38807.1; -.
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                                                                                                                               to the canine genome.";
                                                                                                                                                                                                                               SEQUENCE FROM N.A.
Venta P.J., Gao Y.,
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SEQUENCE FROM N.A.
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Echinoldea: Euechinoldea; Diadematacea; Diadematoida; Diadematidae;
                                                                     MEDLINE-88123478; PubMed-2836084; McCracken A., Uhlenbusch I., Gellissen G.; Structure of the cloned Locusta migratoria mitochondrial genome. Testriction mapping and Sequence of its ND-1(JRF-1) gene.": Curr. Genet. 11:622-530(1987).
EMBL; X05286; GAA28905.1; -.
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"Population structure and speciation in Ungical seas: global
phylogeography of the sea urchin Diadema.";
Evolution 55:955-975(2004).
Pteryjota, Nogiona, Orthoptor idea, Orthoptora, Caelifora,
Acridomorpha, Acridoldea, Acrididae, Oedipodinae, Locusta.
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01-MAR-2002 (TrEMBLrel. 20, Last sequence uplate)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
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                                                                                                                                                                                          8 AA; 1019 MW; FBE3372330464588: 18064;
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Rest Local Similarity
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                                                          SEQUENCE FROM N.A.
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Miterhoodrien.
                           NCBI_TaxID-7004;
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SEQUENCE FROM N.A.
STRAIN+AN414, DA4441, DIAN125, DIAN126, DIAN131, CM2, AND TM4:
STRAIN+AN4414, DA4441, DIAN125, DIAN126, DIAN131, CM2, AND TM4:
MEDINE-21561594; Pubmed-11703875,
MESSICS H.A., Garrido M.J., Kessing B.D.;
"Demographic history of Pladema antillarum, a keystone herbivore on Caribbean reefs";
                                                                                                                                                                                                                        STRAIN-SLAIZI AND GLAIZ4:
BEDELINE-21561594: PubMed-11703875:
Lessics H.A., Garrido M.J., Kessing R.D.;
"Demographic history of Diadema antillarum, a keystone herbingre on
Enlargada, Merazna, Entinodermata, tima merimba, especial Bohimoldea, Eucchinoldea, Diadematacca; Diadematoida; Diadematidae;
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Echinoidea, Euechinoidea, Diadematarea, Diadematoida, Diadematlihue,
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                                                                                                                        Lessios H.A., Kessing B.D., Pearse J.S.; "ropulation structure and speciation in tropical seas: global phylogrography of the sea urchin Diadema";
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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100.0%; Pred. No. 6.7e+05;
live 0; Mismatches 0; Indel
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UMAR-2002 (TrEMBLEL 20, Last sequence update)
ULMAR-2002 (TrEMBLEL 20, Last amortation update)
Cytochrome oxidase subunit II (Fraqment).
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                                                                            SEQUENCE FROM N.A.
STRAIN-GLAIZI, AND GLAI24;
MEDLINE-21323357; Pubmed-11430656;
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Proc. P. Soc Lond
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Best Local Similarity
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                                           NCB1_Tax1D=105359;
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6 LE 7
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STRAIN-CLS. ADD. D6. GG64, G123, DM1. FM - AND DM71:
STRAIN-CLS. MEDLING-21561534 : PubMed-11703875;
Lessios H.A. (darrido M.J. Kessing B.D.;
"Demographic history of Diadema antillatum, a keystone herbivore on
                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota, Metazoa, Echinodermata, Eleuther voa, Echinozoa,
Echinoidea, Poechinoidea, Diadematacea, Dia kwatoida, Diademafidae,
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Evolution 55:955-975(2001).
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STRAIN-C15, D3. D5, D6, CC66, G123, DM1, DM: AND DM71;
MEDLINE-21323357; PubMcd-11430655;
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Proc. R. Soc. Gond., R. Biol. Sci. 268:2347::353(2001).
EMBL. AY012901: AAL33887.1: -...
EMBL. AY012911: AAL33889.1: -...
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              NON_TER 1 1 1 SEQUENCE 8 AA; 1037 MW; 701B173B46DDC2EG GRC64;
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Best Local Similarity 100.0%; Pred. No. 6.7++05;
Matches 2; Conservative 0; Mismatches 0;
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EMBL, AY012919, AAL33842.1; --
EMBL, AY012940; AAL33844.1; --
EMBL, AY012949; AAL33849.1; --
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Matches 2, Consorrative
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01-MAR-2002 (TrEMBLrel.
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ORWBG4;
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SO FT
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Eukaryota, Metazoa, Echinodermata, Eleutherozoa, Echinozoa,
Echinoidea, Fuechinoidea, Diadematacoa, Diadematuidae, Diadematidae.
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100.0%; Pred. w.
Cytochrome oxidase subunit II (Fragment).
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                                                                                     Diadema paucispinum.
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April 29, 2003, 08:30:19 ; Search time 42.0833 Seconds (without alignments) 15.832 Million cell updates/sec
GenCore version 5.1.4 _{\rm L}F_{\rm 2}-4578 Copyright (c) 1993 - 2003 Compagen Ltd.
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Rost-processing: Listing litst 45 summaries state of the summaries of the

Minimum DB seq length: 0 Maximum DB seq length: 10

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size:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	HIV-1 gpl20 induce	Human peptide #152	Vascular dementia-	Human API-148 tryF	DPI tryptic digest	Depression-Associa	Human neurofilamen	Human neurofilamen	Peptide for analys	AIDS virus recepto
SUMMARIES	AAY49695	AAM98253	ABB55971	ABB52194	AAU28456	AAU26102	AAY20715	AAY20656	AAE08624	AAP71550
80	07	C1	22	(1 (1	(1 (1	CI CI	10	19	d	80
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s Query Match	0.0	0.0	0.0	0.0	80.0	0.0	c O	0.0	0.0	0.0
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(ADIM-) ADVANCED INMUNIT INC.

WPT; 1999-613605/54. Pert C, Ruff M;

98US-0080836. 94W0-US07514.

06-APR-1999; 06-APP-1998;

W09951254-A1.

14-0CI-1999.

virus cina p	s simplex	9.5	ptide I f.r	eptide I for	n applithment	eptide T varian	ide I vari	ide I analog	. acidoraldar	4	anti-HRS a	e carly	Human heary chain	CD8 beta chain rid	arker.	ectin 3	Galectin 3 7-termi	ernalisation s	ernalisation	nhibiting	celet GF11	derived	an T 0-11		Human 1012 11010-0	cifficals straigs	The specific		Tat prote	uman G-protein.	Sychology applied
AAP71551 AAPR3009	AAR269	AAR2894 AAR4433	AAR4783	AAR4783	AAR7733	AAR6148	AAR5949	AAR:9899	AAR9064	AAW9031	AAB5218	AAB9951	AAR6749	AAB3756	AAB5566	AAB5750	AAB5791	AAR4165	AAR4165	AAR3189	AAR5121	AAR5612	AAW3946	AAW3946	AAY211	AAW8381	AAW837	SEJ LAWY	AAY051	AAUO438	WY.
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122	41.	15	17	3:1	19	50	21	검	23	্র	25	90	C •	82	Ä	30	31	3.2	(r)	34	35	36	3,7	38	かと	0.7	,-1 *1	: ;	4. 4.	\$ \$	<u>.</u>

AT LIGHMENTS

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HIV i gpiiù induced neuronal vell death inhibiting peptide #1.
       AAY43635 Standard, pertide: 5 AA.
                     'A LAM-2777 (first entry)
                                                              Synthetic.
              AAY49695;
FESULT 1
   AAY49695
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunosuppressive; immunostimulatory, antiintlammatory; cytostatic; enuroportective; antimicrobial; gene therapy; vectorie; amplase; cancer; amyloid protein; andiopotetin; apoptosis related protein; cadherin; cyclin; polymenase; encogene; histone; kinase; colony stimulating factor; copplement related protein; cytochrome; Kinasin; cytokine; interferou; interleukhi; G-protein; coupled receptor; thi, esterase; inflammation; multifactorial disease; autoimmune disease; infection;
                                                                                                                                                                                                                      induced neuronal cell death. Phainaceutical compositions containing the peptides are useful for treating symptoms caused by neuronal cell loss. Such conditions especially associated with HTV infection include encephalogathies, neuropathies, memory loss. Computed, depression psychosis and opportunistic infections. The populars and as annayonists of gpilo-mediated neurotoxicity and subsequent memory legislated bygeneration. This enables therapeutic treatment of HTV infection and other inflammatory neurological diseases, including multiple sclerosis, tropical spastic paraparesis and Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymorphic nucleic acids encoding e.g. ampliases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections
New peptides useful for inhibiting human inmen 3-deficiency virus type 1 (HIV-1) gp120 induced neuronal cell death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                  AAY49695 and AAY49696 represent peptides which inhibit HIV-i gpiż0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human peptide #1528 encoded by a SNP oingonn-leotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 5; DB 2 : I
100.0%; Pred No. 7.8e+95;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 4003; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAM98253 standard; Peptide; 9 AA
                                                                                                                       Claim 1; Page 11; 16pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-DEC-2000; 20008-0173419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100 0
nes 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shimkets RA, Leach M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI: 2001-465210/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200147944-A2.
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1 LESYT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               wery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       itches
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The invention relates to screening, diagnosis or prognosis of Vascular Dementia (VD) in a subject comprising analysing body fluid from the subject to progness to generate a 2-D array of features containing at least one chosen feature whose relative abundance correlates with the presence, absence, stage or severity of VD or predicts the onest or course of VD. especially detecting in a sample of cetebrospinal fluid (CSF) from the subject one of 22 VD-associated protein isoforms (VFIS) (ARRESRO)-ARREA25) as fully defined in the specification. Detecting VD-associated determined in the screening, diagnosis or prognosis of VD, for determining the stage or severity of VD, for identifying a subject at risk of VD or for
complement related proteins, cytochromes, kinesins, cytokines, interleukins, G-protein coupled receptors and thioestelases. Interleucins, interleukins, G-protein coupled receptors and thioestelases. The present sequence is a peptide enroded by one such alignment extraction, diagnosis and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with the prevention, of the proteins listed above. Discribers that may be prevented, diagnosed antitor treated include multifactorial diseases with a genetic component, such as autolimmine diseases (e.g. rheumatoid architis multiple sclerasis, diahetes, systemic lupus erviticematoid architis multiple sclerasis, diahetes, systemic lupus erviticematoid bialic, treast, calor mark fidony, lephacemia), diseases of the performs system and an infection of pathogenic organisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Screening, diagnosis or prognosis of mascular dementia (VL), useful for determining stage of VD and monitoring the effect of VD therapy, comprises analysing body fluid by 2-dimensional electrophoresis for features correlated with VD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vascular Dementia, VD; VD-associated protein isoform; VPI: screeting:
                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                              Score 4: DB 22; Length 9: ; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vascular dementia-associated protein isoform (VPI) 171.
                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (OXFO-) OXFORD GLYCOSCIENCES UK LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diagnosis, prognosis, gene therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABB55971 standard; Peptide; 10 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 6; Page 33; 151pp; English.
                                                                                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14 MAR 2001; 2001WO-GB011(6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-Nov-zagan; zhágats-0724391.
                                                                                                                                                                                                                                                                                                                                    80.08
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                                                                                                                                                                                                                                                                                                                                                                             4: Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Herath BMAC, Farekh RB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WP1, 2601-557937/62.
                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                            9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       W0200169261-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24 - NOV 2000:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                        1 LESY 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 LESY 5
                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB55971;
                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 3
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1 LESY 4
                           3 LESY 6
                                                                                  AAU28456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Screening for Authenmer's disease in a manual, by making two-dimensional array of a feature whose relative abundance correlates with disease, and compating with abundance of the feature in samples of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is a peptide produced from an API by proteolysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disease-Associated Protein Isoforms (APIs) in cerebrospinal fluid, serum or plasma. The abundance of the AFS and APIs is then normalised to an Expression Reference Protein Isoform (BRPI) in order to determine whether a patient is suffering from, or has a predisposition to, Alibeimer's Disease. The relative abundance of the AFS and APIs correlates with the severity of Alzheimer's Disease.
monitoring the effect of therapy administered () a subject having VD. Nucleic acids encoding a VPI or inhibiting the function of a VPI are useful for the treatment of VD and for gene therapy.
                                                                                               Saps
                                                                                                                                                                                                                                                                                                 Human; neuroprotective; nootropic; gene therap;; vaccine;
Alzheimer's disease; Alzheimer's Disease-Assochated Feature; AF;
Alzheimer's Disease-Associated Protein Isoform, API; tryptic digest;
Expression Reference Protein Isoform; ERPI; priveolysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to methods for the screening, diagnosis and proquosis of Alzheimer's disease. The methods involve the detection of Alzheimer's Disease-Associated Features (AFS) and Alzheimer's
                                                                                              ċ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parekh RB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Durham KL, Friedman DL, Herath HMAC, Kimmel LH, Parekh RB Potter DM, Rohlff C, Silber BM, Stiger 1E, Sunderland P1: Townsend RR, White F, Williams SA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; D8 22; Leus
No. 136+12;
9; Indels
                                                                       Length 10;
                                                                                            e; Indels
                                                                      DB 22; Ler
. 1.3e+0.;
                                                                      Score 4; DB 22; Pred. No. 1.3e
                                                                                                                                                                                                                                                                           Human API-148 tryptic digest peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (OXFO-) OXFORD GLYCOSCIENCES UK LID
(PFIZ ) PFIZER INC.
                                                                                                                                                                                                      ABB52194 standard; Feptide; 10 AA.
                                                             80.0%; Scur.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example: Paye 30; 162pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-APR-2000; 2000US-194504P
28-NOV-2000; 2000US-253647P.
                                                                                                                                                                                                                                                                                                                                                                                                                                3-APR-2001; 2001Wn-0510908
                                                                                                                                                                                                                                                    08-FEB-2002 (flrst entry)
                                                                                 Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rohlff C,
White F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-679484/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          healthy persons
                                                                                                                                                                                                                                                                                                                                                                                   W0200175454-A2
                                                                                                                                                                                                                                                                                                                                                              HOMO Sapiens
                                                                                                                                  3 LESY 6
                                                                                                                     1 LESY 4
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                                                  Seguence
                                                                        Query Match
                                                                                               Matches
                                                                                                                                                                               RESULT 4
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80.0%; Score 4; DB 2 100.0%; Pred. No. 1.3 tive 0; Mismatcles

Query Match Best Local Similarity 100. Matches 4; Conservative

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Novel nucleic acid encoding a protein associated with bipolar affective disorder, which is used for diagnosis, prophylaxis and therapy of neuropsychiatric disorders, such as bipolar affective disorder.
                                                                                                                                                                                                                                                                  Human: depression associated protein isoform: tryptic didest peptide:
DPT, cerebouspinal fluid, CSF, BAD, bipolar affective disorder:
neuropsychiatric disorder: bipolar mood disorder; neuroleptic;
maniac-depressive illness: schizoaffective disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to the identification of depression associated protein isoforms (PPIs), particularly the tryptic digest peptides of these proteins. Some of the PPIs (AAU28404-AAU28625) described are decreased in the merebrospinal fluid (ASF) of FAL (Dipolar affective disorder) subjects, whilst other PPIs (AAU28626-AAU28627) are increased in PAP subjects. Also described are peptide Sequences identified from DPI-45 and PPI-213 and the nucleic acid sequence they are encoded by. The sequences of the invention are useful for chinical screening, diagnosis, prognesis, therapy and prophylaxis of neuropsychiatric disorders e.g. BAD (also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             attention deficit disorders, schizoaffective disorders, and unitariar affective disorders. The present sequence represents one of the IFI tryptic digest peptides of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   known as bigolar mood disorder. 89), maniac-depressive illnesses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DR 22; Length 10; 1.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Herath HMAC, Parekh RB, Rohlif C. Terrett JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 4; DR 22; Pred. No. 1.3e (): Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (OXFO-) OXFORD GLYCOSCIENCES HK LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure, Page 31, 153pp, English.
AAU28456 standard: Peptide: 10 AA
                                                                                                                                                                                                          DPI tryptic digest peptide #53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.0%; ses 4, Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-FEB-2000; 2000GB-0004412.
08-086-2000; 2000GR-0030050
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-DEC-2000; 2000US-0254830.
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                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WC206162787-A1.
                                                                                                                                        03-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 LESY 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3 LESY
                                                              AAU28456;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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AAU26102
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22-TTT-1999 (tilst Autiv)
  The invention relates to a preparation composing an isolated Bipolar Affected Disorder (BAD)-Associated Protein isoform (DPIs). The DPI's are used to screen, diagnose or prognose of BAD or unipolar depression, determine the stage or severity of BAD or unipolar depression, identify a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         attention deficient disorder, a schizoaffective disorder, a hipplan or a unipolar affective disorder. The bris are uned in proteomics. The proteomic approach of using DPIS for screening, diagnosis or prognosis of BAD or unipolar depression overcomes the proteoms of using gene expression analysis, such as not being able to obtain central nervous system (CNS) tissue from a living patient under normal circumstances. The present sequence is a DIP decreased in the CSF (cerebro-spinal fluid) of subjects having BAD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      subject at risk of developing BAD or unipolar depression, or modified the effect of therapy in a subject. They are also used to screen for or identify agents that interact with a DPI. Those agents, antibodies against the DPIs, and nucleic acids encoding the DPIs are used to treat or prevent BAD or unipolar depression. Diseases that can be treated are
                                                                                                                                                                                                                                                                                                                                                                                                                          Preparation for diaynosing or treating bipciar affected disorder (BAD) or unipolar depression, or for screening for modulators, comprises a BAD-associated protein isoform .
                                                                                                     Human, Bipolar Affective Disorder, BAD, Depression-Associated Contact
DF: Depression-Associated protein isoform, Fig. Devetro-Spinal fluid,
GSF: antidepressant: antimanic; nootropic: Usnguiliser: neuroleptic:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                              attention deficient disorder; schizoaffective disorder:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        80.0%, Scare 4, Jalum 1. 190.0%; Pred No. 1.344.32;
                                                                               Depression-Associated Protein isoform DPI-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                              (OXFO+) GXP(+PD GLYCSCIENCES UK 11P
                                                                                                                                                                                                                                                                                                                                                                        Herath HMAC, Parekh RB, Rohlff C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY20715 standard; Protein; 4 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 8; Page 31; 163pp; English.
  AAU26102 standard: Peptide: 10 AA
                                                                                                                                                                                                                                                                 23-FEB-2001; 2J01WO-GB00791.
                                                                                                                                                                                                                                                                                                         200068-0030050.
                                                                                                                                                                                                                                                                                            24-FEB-2000; 2300GB-0004412.
                                                                                                                                                                                                                                                                                                                     200008-0254830.
                                                                                                                                                           anipolar attective disorder
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                                                      (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 AA:
                                                                                                                                                                                                              W0200163294-A2.
                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                        38-DEC-2000;
12-DEC-2000;
                                                    18-DEC-2001
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3 LESY 6
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  CCCCCCXXXHHHXXXXXXXXXXXXXX
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This invention describes a novel method for the diagnosis of a disease caused by, or associated with, an RNA molecule that has a transmitt mutation. The method is used to diagnose age-related diseases, especially caused by and a wide range of diseases, especially disease, bown's syndiome, myotonic district (e.g. Alzhenmen's disease, bown's syndiome, myotonic district (is ease mithing the Alzhenmen's disease, allows a definitive diagnosis of Alzhenmen's disease in living patients, allows a definitive diagnosis of Alzhenmen's disease in living patients, and meanly stage. It is based on the observation that disease may be caused by mutations in RNA makerules, secifically proteins including beta amyloid precursor protein (beta-APP), the microtubule associated protein 2 (MAP2), neurofilament—E, neurofilament—M, neurofilament—E, neurofilament—E, neurofilament—E, neurofilament—E, neurofilament—E, neurofilament—E, neurofilament—E, transperient (AR2), presenting II, gital fibrillary action of protein (AR2), neurofilament—E, neurofilament—E, transperient (AR2), presenting III, HEFF-1, High m. Lilly protein A. England C. (Edic.) If the mid-open contribution of the control of the cont
                                                                       Saps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Diagnosing discase ky detecting frameshift mutations in PNA or corresponding protein mutations - used to diagnose cancer and neurological diseases, particularly Alzhelmer's disease, and also for treatment and prevention with specific ribozymes or wild-type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indeis
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Human neurofilament-M wild type protein fragment 57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     60.0%; Score 3; DB 19; Le
100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Burbach JPH, Grosveld FG, Van Leeuwen FW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HOYA-) HOYAL NETHERLANDS ACAD ARTS & SCI.
(UYRO-) UNIV POTTERDAM ERASMUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Figure 8: 258pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N-PSDB; AAX75759.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-APR-1998;
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AAY20656

RESULT 9

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This invention describes a nevel method for the diagnosis of a disease caused by, or associated with, an RNA moternal that has a frameshift mutation. The method is used to diagnose age related diseases, especially cancer and a wile rarge of neurogegenerality Huntington's disease, in disease, Down's syndrome, myotonic dystrophy, Huntington's disease, allows a defenditive diagnosis of alsease, disease mellium type II and many others listed) or susceptibility to these disorders. The method allows a definitive diagnosis of Alzheimer's disease in living patients, at an early stage. It is based on the observation that disease may be caused by mutations in RNA rather than DNA. The invention describes the used of neuronal system RNA motecules, specifically proteins including porteins from precursor protein (beta APP), the microtubule associated protein 2 (MAP2), neurofilament. In microtubule associated protein 2 (MAP2), neurofilament. In quial fibrillary acidic neurofilament. Protein (GFAP), the ceilaid Lumbur antigen protein and but the ceilaid Lumbur antigen protein (GFAP), the ceilaid Lumbur antigen protein protein protein antigen protein antigen protein pro
                                                                                                                                                                                                                                                                                          Alzheimer's disease, 194 interpretation and agentifications. Alzheimer's disease, multiple scherosis, alcouble dystrophy, neuronal, Huntington's disease, multiple scherosis, alcouble strophy, neuronal, disease, multiple scherosis, alcouble sold protecting fau, big Tau, ubdgattin B, apoilepercieto E, MAR2; neurofilament-E; neurofilament-E; neurofilament-E; presentlin I; rellular tumour antigen, alla [Lukhana and added to protection of psis semaphorin II; Hurb-I; bell leukemia/lymphona 2 protectionogy-re; HWRP-C; NSP-A; high mobility group protein C, neuroendocrine specific protein A.
                                                                                                                                                                                                                                                                       age-related disease; neur-degenerative disorder;
                                                                                                                                                                                                                                           Human, beta-amyloid precursor protein, beta-Aff; diagnosis; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           corresponding protein mutations - used to diamnose cancer and neurological diseases, particularly Alzheimer's disease, and also for treatment and prevention with specific of agrees or wild-type.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diagnosing disease by detecting frameshift mulations in ENA or
                                                                                                                                                                            Human neurofilament-L wild type protein fragment 46.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Grosveld FG, Van Leeuwen FW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ROYA-) ROYAL NETHERLANDS ACAD ARTS & SCI
(UYRO-) UNIV ROTTERDAM ERASMUS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure: Figure 7: 258pp: English.
AAY20656 standard; Protein; 4 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98W0-IB00705
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                                                                                                                   22-JUL-1999 (first entry)
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                                                                                                                                                                                                                                                                       frameshift mutation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI: 1998-60990'/51.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                              AAY20656;
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gamma induction activities. The polypeptides and polynucleofides of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention are useful for treating various diseases e.g. bacterial infections. Albs, diabetic retinopathy and cancer. The polypeptide and perlymorleotide may also be useful as a vaccine adjuvant, to enhance a vaccinated host's cell-mediated immunity for protective response to a pathogen. The present sequence is a peptide used in the study of Troell proliferative activity of human cytokine P35/P40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to nucleic acids encoding modified stacking polymeptides (1.0. modified p40 and p35 polymeptides). The modified cytoking polymeptides have Theold proliferative and interferon (IFN)-
                                                                                                Peptide for analysica Thopil proliferative antivity of Tuman p35/P40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Muchelo acids encoding modified cytching polypeptides (i.e. modified p40 and p45 polypeptides), useful for treating various discases e.u. diabetic retinopathy and cancer
                                                                                                                                                   AIDS: diabetic retinopathy; cancer; vaccine; cell-mediated immunity;
                                                                                                                                 Tytoking: Tropli: interferon-gamma: IFN gamma: bacterial inferion-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60.0%; Score 3; PB 22; Le
Liarity 100.g%; Fred. Mu. 7.8e+05;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AIDS virus receptor site blocking peptide (III).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bisclusure: Page 31: 223pp: English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAP71550 standard, Protein; 5 AA.
AAE08624 standard; peptide; 4 AA
                                                                                                                                                                                                                                                                                                    30-NOV-2000; 2000WO-US32664.
                                                                                                                                                                                                                                                                                                                                      99015-0169035
                                                                                                                                                                                                                                                                                                                                                       28-NOV-2000; 2000US-0169035
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                                                                01-NOV-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                      Leong sk, Punnonen J:
                                                                                                                                                                                                                                                                                                                                                                                        (MAXY-) MAXYGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-502381/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          focal Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 AA;
                                                                                                                                                                                                                                     W0200140257-A2.
                                                                                                                                                                                                                                                                                                                                      02-DEC-1999;
                                                                                                                                                                                                                                                                     07 - JUN - 2001.
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                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                      №35; p40.
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Gaps

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Length 4; O; Indels

DB 14; Le . 7.8e+05;

60.0%; Score 3; DB 1 ilarity 100.0%; Pred. No. 7.8 Conservative 0; Mismatches

Local Similarity es 3; Conservat

Query Match Matches

Sequence 4 AA;

9

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87US-0048148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-AUG-1986;
                                   12-DEC-1986;
03-JUN-1987;
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                         26-JUN-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAP83009;
                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                        virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                             quencic formula.

The peptide blocks effectively receptor sites of cells and prevents cell infectivity with AIDS virus in monkey, rated and human brain membranes and cells of the lumnune system.

Vaccine preprs, contg. the peptide provide protection against infection by AIDS virus. The peptide can also be used in Kits for the detection of the AIDS virus and antibodies to the AIDS virus; and as immunogens to elicit monoclonal antibodies to the AIDS virus; the peptide may be produced by conventional antibodies synthesis using solid phase or liquid phase in thouse see also AAP71548-56, and EP-249394.
                                                                                                                                                                                                                                                                                       New peptide(s) related to AIDS virus - which inhibit AIDS virus binding to receptor sites and prevent cell infectivity with AIDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      60.0%; Score 3; DB 8; Length 5; 100.0%; Fred. No. 7,80**5;
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 varine, adibadies.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAP71551 standard; Protein; 5 AA.
 Constitution of the Salve,
                                                                                                                                                                                                                                                                                                                                        Claim 5; Page 9; 12pp; English.
                                                                                                                                                                                                                                         Pert CB, Ruff MR, Farrar WL.
                                                                                                                                      86US-0869919.
86US-0878586.
86US-0940919.
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(USDC ) US SEC OF COMMERCE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AA;
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   Aifes: wirms.
                                                                                                  03-JUN-1987;
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26-JUN-1986;
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                                                                                                                           11-MAY-1987;
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                          Synthetic.
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                                                                                                                                                                                                                                                                                                                   virus
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The peptide blocks effectively receptor sites of cells and prevents cell infectivity with ADDS virus in monkey, rat and human brain membranes and cells of the immune system. Vaccine prepris contg the peptide provide protection against infection by ADDS virus. The peptide can also be used in kits for the detection of the ADDS virus and antibodies to the ADDS virus: and as immunogens to elicit monoclonal antibodies. The peptide may be produced by conventional methods of peptide synthesis using solid phase or liquid phase methods.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Blocking peptide used in composition for treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                         New peptide(s) related to AIDS wirus - which inhibit AIDS wirus binding to receptor sites and prevent cell infectivity with AIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monotional antibodies and peptide(s) - useful for treatment and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This peptide is a specifically claimed example of a highly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60.0%; Score 3: DB 8: Length 5:
100 0%; Pred No 7 8e+05;
Live 0: Mismatches 0: Indels
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100 08; Pre-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           claim 5; Page 9; 12pp; English.
                                                                                                                                                                                                                                                       Pert CB, Ruff MR, Farrar WL;
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8605-0898273.
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                                                               86US-0940919.
                                                                                          87EP-0304913.
86US-0869919.
                          86US-0878586
                                                                                                                                                      (PERT/) PERT C B. (USDC ) US SEC OF COMMERCE.
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has hever been reported to penetrate any type of cells, but only to occupy the CD4 receptors found on lymphocytes and it apparently penetrates the stan with relative ease. A major advantage of the penetrates the stan with relative ease, to take the standard of the penetrates the stan with the last of toxicity, which means that the nay its days for an extended period of time is possible without any itamines. It is an extended period of time is possible without whicheves I it is an extended a highly generic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The peptide is useful in the treatment and prevention of diseases or conditions caused by Helpes viruses, esp. Helpes Simplex Virus 1 (HEV I), HSV-2, Validella Simplex Virus 1 (HEV) or Epstein Teat Value (EBV). It can also be used to treat diseases and conditions caused by the family of helpes viruses in animals, esp. Adjessivy's disease in pins, boother innoitable list in helpes and properties in production to the period disease in chickens. It can also be used in dispnostics, the period
                                                                                                                                                                                                                                                                                                                                                                                            HSV; HSV-1; HSV-2; Varicella Zester virus; human cytomegalovirus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       occupy lymphocyte CD4 relations for theating and preventing virus infections with Harpes simpley. Varioella Zoster, cytomegalovirus and Epstein-Barr virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New linear or cyclic peptide(s) including cysteine residue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             O: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DR 13; Length 5;
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  Mismatches
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                 Herpes simplex virus treatment peptide.
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                                                                                                                                                                                                          AAP26903 standard, peptide; 5 AA
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  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                    Epstein-Barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
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                                             3 SYT 5
                                                                                      3 SYT 5
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  Matches
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                                                                                                                                                                                  AAR26903
                                                                                                                                                                                                                                                          This peptide is used in a composition for treatment of thronic conditions of neuropsychiatric disorders and psoriasis through inhibition of HIV cell surface receptor (cb4) binding. Cb4 site is common to the CNS and immune system. AIDS- and non AIDS-related psoriasis both respond favourably to treatment with the peptide-containing composition. There are a range of forms which the composition and administration can take.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US Government-cowned NTIS applications to prevent classes with ongoing ^{13} granted patent numbers. For further information please visit the Derwent web site at www.derwent com/dwpi/updates/ntis us html )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The serine residue can be replaced by an asparagine residue. Analogues of these peptides with D-Threonine as the amino-terminal residue and/or an amide derivate at the expectations are also be used. (Note: Revised entry submitted to correct the patent number format of
                                                                                    This is a blocking peptide which inhibits virus proliferation. It is useful, in conjunction with a monoclona, antibody to a neutralising region of HIV, for treatment or prophylaxis of HIV infections. N. and C-terminals can have up to 20 amino acids attached. See also AAPB3003-08 and AAPB3010-12.
                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pentapeptide Which inhibits HIV binding to cell surface receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Use of short pertide(s) to inhibit binding of HIV to human cells treating psoriasis and neuropsychlatric discrete including memory
                                                                                                                                                                                                            This is equivalent to NL8701950 (accession no is from this)
                                                                                                                                                                                                                                                                                                                                                 : Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60.0%; Score 3; DR 10; Length 5; 160.0%; Pred. No. 7.8e-v.5,
                                                                                                                                                                                                                                                                                                   DB 9; length 5;
  diagnosis of human immuno:deficiency virus infections
                                                                                                                                                                                                                                                                                                            100.0%; Pred. ....
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                                                                                                                                                                                                                                                                                                60.0%; Score 3; L
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (USSH ) US DEPT HEALTH AND HUMAN.
                                           Claim 42; Page 24; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAP91855 standard; protein; 5 AA.
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Best Local Similarity Tuo.v
And 3, Conservative
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AAP91855;

RESULT 13 AAP91855

Sednence

Sequence

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                      CES: Latigue; tension; anger; confusion; poptide T; HIV; synthetic; human immunodeficiency virus; gp 120.
                                                            Example of a generic peptide for treating chionic fatique syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 60.08; Score 3: DB 1: Length 5: Best Local Similarity 100.08; Pred. No. 7.8e \cdot 05; Matches 3; Conservative 0; Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptide compsns, for treating chronic fations syndrome ameliorate symptoms and improve vigour and sognitive and neuro-motor performance
                                                                                                                                                                                                                /note= "may be amidated"
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                                                                                                                                                  Location/Qualitiers
Misc-difference 1
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                                       23-MAR-1993 (first entry)
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                                                                                                                                                                                                      Modified-site
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                                                                                                                                   Synthetic.
             AAR28941;
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GenCore version 5.1.4_05_4578 Coffyright (c) 1993 - 2003 Compound Ltd.
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Searth orme in Searnds (without alignments) 40.065 Million cell updates/sec April 29, 2004, 98 52 00 Run on.

50 duence: 1 LESYT 5

Sequence

Gapop 60.0 , Gapext 60.0 Scoring table, oligo

saula seys, Anl? 9803 residues Seagrehad.

Frd Size :

al number of hits satisfying chosen parametern

Minimum DB seq length: Û Maximum DB seq length: 10

//guz_//ptodata/Z/pubpaa/TSOO_pubchamb_pop. Publ_/_prodata/Z/pubpaa/TSOO_pubchamb_pop. //guz_S/prodata/Z/pubpaa/TSOO_pubchamb_pop. //guz_S/ptodata/Z/pubpaa/TSOO_pubchambub.pop. FILES JAK. FOR. * . gan. I., predataki, pubbaa, horro, pubbi.MR pop. *//ogn2_6/prodataki/pubpaa,⁴¹⁸⁰⁹_FOBGOMB pop. *//ogn2_6/prodataki/pubpaa, 3500, REM_FOB. pep. */

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Score		, n Length	DB	ID	Description
4	0.08	10	6	US-09-825-290 200	Sequence 209. Apr
-+	0.09	07	۳,	3 - 536 - 878 - 50 - 55	Sequence 6. Applil
4	80.0	10	9	US-09-791-389-4".	
4	0.08	10	Œ.	#8-00-201-301-4v	
m	90.0	T	٦.	715 - 05 d - 120 - 50 - SH	-T
3	11 11	u -	- 1 - 1	1.7 · 0.47 · 987 · 50 - 50	Seditebre 200 April
1,50	0 09	,- L	€.	11.4-04-828-80-811	
m	U ()	۸	σ	4-8150-659-00-SII	Ā
æ	0.09	J 0	2,	08-09-995-775-4	Sequence 47, Appl
m	0.09	9 0	σı	US-10-097-175-99	Sequence 99, Appl
e	0.09	9 (10	US-09-984-292-40	Sequence 40, Appl
e	50.09	9	10	1 4-264-686-60-30	Sequence 40, Appl
~	9.09) 6	10	US-09-867-852-1:8	Sequence 138, App
κ.	0.09	7	6	US-10-000-273-1	Sequence 13, Appl
3	0.09) 7	ı, Di	US-10-091-246-15	Sequence 15, Appl
٣	60.0	7	6	US-09-879-461-18	Sequence 18, Appl
m	U 09	7	σ	US-09-884-7678-, (0)	100,
3	0.09) 7	Þ	US-09-977-797A-13	Sequence 38, Appl
₹.	60.0) 7	.5.	08-10-091-313-5	
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i LESY 4

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ALTGNMENTS

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APPLICANT: Williams Stephen A.
TITLE OF INVENTION. Nucleic Add Molecules, Polypeptides and
TITLE OF INVENTION. Uses Therefor, Including Diagnosis and Treatment of
TITLE OF INVENTION: Alzheimer's Disease
FILE REFERENCE, 2572-1-021 NZ
CORRENT AFFLICATION NUMBER, US,CS,925,230
CORRENT AFFLICATION NUMBER, US,CS,925,230
FRICE AFFLICATION NUMBER, US, CS,194,504
PRIOR APPLICATION NUMBER, NS,CS,93,647
PRIOR APPLICATION NUMBER, NS,CS,93,647
PRIOR PILING DATE: 2000-11-28
NUMBER OF SEQ ID POS WINDOWS Vergion 4.0
SEQ ID POS 999
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100.0%; Pred. No. 43;
cive 0; Mismatches
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White, Frost
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Hest Local Similarity 100.0
These 4: Conservative
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ORGANISM: homo sapien
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RESULT 1
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us-09-647-749a-1.oliszlm.rapb

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(415) 576-0200
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ORGANISM: homo sapien
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Matches 4: Conservat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ns-00-791-393-46
US-09-791-393-46
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STATE:
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APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Parekh, Rajesh Bhikhu
APPLICANT: Terrett, Jonathan Alexander
APPLICANT: Terrett, Jonathan Alexander
APPLICANT: Trent Louise
TITLE OF INVENTION: Proteins, Genes and Their Use for
TITLE OF INVENTION: and Unipolar Depression
TITLE OF INVENTION: and Unipolar Depression
FILE REPRENCE: 2343-1-001 N.2
UPPENT FILING DATE: 2001-22-3
PRIOR APPLICATION NUMBER: GB 0044412-3
PRIOR FILING DATE: 2000-12-08
PRIOR FILING DATE: 2000-12-08
PRIOR FILING DATE: 2000-12-12
PRIOR FILING DATE: 2000-12-12
PRIOR FILING DATE: 2000-12-12
PRIOR FILING DATE: 2000-12-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    80.0%; Score 4; DR 9; Length 10; 100.0%; Pred. No. 43;
                                                                                                                                                                                                                                                                                                             APPLICANT: CALENOFE, EMANUEL.
APPLICANT: CALENOFE, CHARLES C.
TITLE C. INVENTION: IMMUNISENIC PEPTITIES AND THE STELLS FILE REPERENCE: 14417, 2114 (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (114) (11
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Best Local Similarity 100.0%; Pred. No. 433
Matches 4; Conservative 0; Mismatches
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Publication No. US20030032773A1
GENERAL INFORMATION:
                                                                                                                                                                                                Sequence 6, Application Hg/09948967
Publication No. US20020193295Al
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Helicobacter pylori
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LENGTH: 10
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Sequence 46. Application #S/09791393
Publication No. US20330312200A1
GENERAL INFORMATION:
APPLICANT: Herath, Herath Mudiyanselage Athula Chandrasiri
APPLICANT: Parekh, Fajesh Bhikhu
APPLICANT: Rohlff, Christian
TITLE OF INVENTION: Proteins, Genes and Their Use for
ITLE OF INVENTION: Diagnosis and Treathment of Bipolar Affective Discussion
FILE PEPERNOR: 2543-1-001 N1
"HEERIT AFFLI WINRES. 2532-01-01 N1
"HEERIT AFFLI WINRES. 2532-01-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Goldberg, Robert B.
TITLE OF INVENTION: Nucleic Acids That Control Seed and
TITLE OF INVENTION: Fruit Development in Plants
NUMBER OF SECUENCES: 324
CORPESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 9; Length 10;
5, 43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Townsend and Townsend and Crew LLP
Two Embarcaders Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     $11001380-070850
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100.0%; Pred. No. 43;
ative 0; Mismatches
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EARLIER APPLICATION NUMBER: GR 0004412.3
EARLIER FILLING DATE, 2000-02 24
EARLIER APPLICATION NUMBER: GB 0030050.9
EARLIER FILING DATE, 2000-12-08
EARLIER FILING DATE, 2000-12-08
EARLIER FILING DATE, 2000-12-08
EARLIER FILING DATE, 2000-12-08
EARLIER FILING DATE, 2000-12-12
NUMBER OF SEQ ID NOS: 308
SOFTHARE: FASLSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 247, Application CS/09071838
Fatent No. US20020152501A1
GENERAL INFORMATION:
APPLICANT: Fischer, Robert L.
APPLICANT: Ohad, Nir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34,774
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IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Yadegari, Ramin
APPLICANT: Margossian, Linda
APPLICANT: Harada, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kiyosue, Tomohiro
Yadegail, Ramin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ALTOFNET/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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0: Indels
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                                                                                                                                                                                                                                                                                                                                                              NAME, RET. DOMAIN LOCATION: (1) ...(6) STREE INFERMATE H. H. FOR STAIN ELLENXXL MODIF
                                                                       Sequence 8, Application US/09953031A
Patent No. US20020177177A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              // Sequence 47, Application US/09995225
// Publication No. US20020193584A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM. Home supjens
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                HS-09-953-031A-8
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                                                            US-09-953-031A-8
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                                                                                                                                                                                                                                                                                        SEO ID NO 8
LENGIH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 9
                                               RESULT 8
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APPLICANT: Burton, D.
APPLICANT: Burton, D.
APPLICANT: Burton, D.
APPLICANT: Schalter, M.
ITLE OF INVENTION: Automnubedies to glucose P. phosphate isomerase and their particitive of INVENTICAL automnume disease
FILE REFERENCE: 1861.005051
FILE REFERENCE: 1861.005051
CURRENT APPLICATION NOMBER: 251.0574
NUMBER OF SEQ ID NOS. 123
SOFTRARE: FastSEQ for Windows Version 4.0
SEQ ID NO. 17
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TITLE OF INVERTION. Chimeric and Humanized Antil Jies to Anglogonin
FILE REFERENCE: 10498/74073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 60.0%, Store 3, CB 11 Length 5, Best Local Similarity 100.0%, Pred. No. 2.76-55, Matches 3; Conservative 0; Mismatches 0, Indels
                                                                                                                                     60 0%; Score %; DR 10; Length 4; 100.0%, Pred. No. 2.7e(5); Liddls tive 3, Mismatches indels
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60.0%; Score 3; DB 1:: Le
Rest Local Similarity 100.0%; Pred. No. 2.7****;
Matches 3; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLI VALUE NO. 256, 242 CURRENT FILING DATE: 1999-04-05 SOFTWARE OF SED ID NOS: 25 SOFTWARE Patentli, Ver. 2.0 SED ID NO. 20 LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 17, Application US, Stries 12 tent No. US2002014675341 JAERAL INFORMATION:
                                                                                                                                                                                                                                                                                                       Paguence 20, Application US/09286240 tent No. US20020(10320A)
TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 2 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                    Matches 3, Jouseirative
                                               4 amino acids
                                                                                         MOLECULE TYPE: peptide US-09-071-838-247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Mus musculus
                                                         TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                       tent No. US2002001
NERAL INFORMATION.
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US-99-286-240-20
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US-09-828-708-17
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TITLE OF INVENTION: G-ALPHA-Q PROTEIN VARIANT AND THEIP USE IN THE
TITLE OF INVENTION: GANALYSIS AND DISCOVERY OF AGONISTS AND ANTAGONISTS OF
TITLE OF INVENTION: CHEMOSENSORY RECEPTORS
FILE REFERENCE: 078003-0280649
                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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APPLICANT: MUBLIER, JOHN L.
APPLICANT: MUBLIER, JOHN
APPLICANT: OZA, VUBHA B.
APPLICANT: COZA, VUBHA B.
APPLICANT: FINDEIS, MAPK A.
TITLE OF INVENITON: PEPTIDIC MODULATOPS OF THE ANDPOSEN PEPEPIOP
CUPPENT APPLICATION UNMER: US/10/097,175
CUPPENT APPLICATION NUMBER: GA/275,240
PRIOR APPLICATION NUMBER: 66/275,240
PRIOR PILING JATE: 2002-01-28
NUMBER OF SEO ID NOS: 102
SOFTHARE: PASISEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Androgen Receptor Finding onlypeptides 10-097-175-99
                                                                                                                                                                                                                                                                                                                                                60.0%; Score 3; DB 9; Length 6; 100.0%; Pred No. 2,76:05; tive 0; Mismatches 0; Indel.
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PELSE APPLITATION NUMBER 10,201,117
PRIOR FILLING DATE: 2001-05-14
PRIOR APPLICATION NUMBER: 50,700,203
PRIOR FILLING DATE: 2001-07-31
NUMBER OF SEQ ID NOS: 67
SOFTWARE: Patentin version 3.1
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PRIOR APPLICATION NUMBER: 60/243,770
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Publication No. US20030045680A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 40, Application 72,79484232
Patent No. US20020128433A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                           ORGANISM Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 3; Conservative
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Best Local Similarity
Matches 3; Conserv
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APPLICANT: YAO, YONG
APPLICANT: XU, HONG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-10-097-175-99
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2 LES 4
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                                                                                                                                                 SEQ ID NO 47
                                                                                                                                                                                                    TYPE: PRT
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APPLICANT: YAO, YONG APPLICANT: YAO, HONG TITLE OF INVENTION: 3 ALFHA-2 FFOIEIN VARIANTS AND THEIR OF INVENTION: 3 ALFHA-2 FFOIEIN VARIANTS AND THEIR OF INVENTION: CHEMOSENSORY RECEPTORS
TITLE OF INVENTION: CHEMOSENSORY RECEPTORS
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: Illustrative OTHER INFORMATION: N-terminal peptide sequence
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APPLICANT: Staskawicz, Brian J.
APPLICANT: Staskawicz, Brian J.
APPLICANT: Brent, Andrew F.
APPLICANT: Cathleck, Couglas
APPLICANT: Kunkel, Fumiaki
APPLICANT: Kunkel, Balkara N.
APPLICANT: Middinos, Michael N.
APPLICANT: M. J. D.-Liang
TITLE OF INVENTION: PREZ GENE FAMILY, PRIMERE, PROBES, AND
ITLE OF LINVENTION: DETECTION METHODS
                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 60.0%: Score 3; DB 10; Length 6: Similarity 100.0%; Pred. No. 2.7e+05; 3; Conservative 6. Mismatches 0, Indels
                                                                                                                                                                                                                                                                                                                                                Query Match 6.0 0%, Score 3; DB 10, Length 6; Best Local Similarity 100.0%; Pred. No. 2.7e+05; Matches 3; Conservative 0; Mismatches 0; Indels
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PILE PEPEPENCE: 07803-0280735
PURRENT APPLICATION NUMBER- 05/03/089,497
CURRENT FILING DATE: 2001-11-21
FRIPP APPLICATION NUMBER- 03/0484,292
FRIPP FILING SATE: 2001-10-29
FRIPP FILING SATE: 2001-10-29
FRIPP FILING SATE: 2001-10-30
NUMBER- 0F SEC IN NOS: 45
SCFTWARE: Fatentin Ver: 2.1
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Fatont No. USLOSLUI9752481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 40, Application US/09989497; Patent No. US20020143151Al
; GENERAL INFORMATION:
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                                                                                                                                                                           ORGANISM: Artificial Sequence
FFISE FILLES LATE, 100, 17-3
NUMBER OF SEC 1D NOS: 42
SOFTWARE PATENTIN VEY, 2.1
SEC ID NO 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Matches 3; Conserva
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US-09-867-852-138
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3 LES 5
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                                                                                                                                                   TYPE: PRT
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FILE REFERENCE: 00746/254002
CURRENT APPLICATION NUMBER: US.09/867,852
CURRENT FILING DATE: 2001-05-29
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/301,085
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28
PRIOR FILING DATE: EARLIER FILING DATE: 1999-04-28
PRIOR FILING DATE: EARLIER FILING TATE: 1994-09-22
PRIOR FILING DATE: EARLIER FILING DATE: 1994-03-32
PRIOR FILING DATE: EARLIER FILING DATE: 1994-03-32
NUMBER: OF SEC ID NOS: 208
SOFTWARE: FASINGO FOR WINDOWS VERSION 4.0
SEQ ID NO 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Grandearl, Andrew David John
TITLE OF INVENTION, NOVEL SENES ENTLAND LAMASTORIER LINE
MOLECULES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        : Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 60.08_{\odot} Score 3: DE 10: Length 6: Best Local Similarity 100.08_{\odot} Prod No 2.756\pm 0.08 Matches 3: Conservative 0: Mismatches : Indels
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SOFTWARE: FastSEQ for Windows Versia: 2.0b
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
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TELECOMMUTATION: INFORMATION:
TELEPHONE: 617/542-6070
TELEPHAX: 617/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/964,127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/10/000,273
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: SEQUENCE DESCRIPTION: SEQ 1D NO: 13:
US-10-000-273-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: 2rews, Ph.D., L. Lee
REGISTRATION NUMBER: P-43,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 06-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 13, Application US/10000273
Patent No. US20020160386A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 7 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                       ; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
US-09-867-852-138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 200154
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIF: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 14
US-10-000-273-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 LES 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 LES 3
                                                                                                                                                                                                                                                                                                                                LENGTH: 6
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0

Gaps

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0; Indels

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0; Indels
                                                                                                                                                                                                                                                                                       60.0%; Score 3; DB 9; Length 7;
100.0%; Pred. No. 2.7e+05;
t.... v. Hismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                              Search completed: April 29, 2003, 08:55:31
                                                                : Sequence 15, Application us/10091236
; Patent No. US20020168360A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                  Owery Match
Best Local Similarity 100.0
                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Mus sp.
                                                                                                                                                                                                                                                                                                                                                                                        Job time : 10 secs
                                               RESULT 15
TS-10 091:236-15
                                                                                                                                                                                                                                                                                                                                        5 LES 7
          111
1 LES 3
                                                                                                                                                                                                                                                                    US-10-091-236-15
                                                                                                                                                                                                                                                                                                                              1 LES 3
1 LES 3
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GenCore version 5.1.4_P5_4578
Copyright (c) 1993 - 2003 Comparen Ltd.
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April 29, 2003, 08:50:15; Search time 15:4167 Seconds Run on:

(without alignments) 9.543 Million cell updates/sec

V 175 105 154 717 45 No. 1

1 LESYT 5 score: Sequence:

Gapop 60.0 , Gapext 60.0 CLIGO Scoring table:

262574 seqs, 29422922 residues Searched:

, size :

local number of hits satisfying chosen parameters

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

***Control of the control of the con

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	% Query Match Length	E E	QΙ	Description
;		60.09	4	. 7	US-09-177-249-247	Sequence 247, App
2	3	0.09	'n	_	US-08-385-443-5	Sequence 5, Appli
3	3	0.09	S	-	-08-456-	36,
4	3	0.09	S	-	-170-	11,
S	~	60.0	S	-	US-08-170-360-14	14,
9	6	0.09	S	~	US-08-302-83-8	9
7	m	60.0	5	٦	.08-481-84	Sequence 6, Appli
œ	m	60.0	S	<u></u>	(1	36,
æ	m	60.09	5	~	-08-403-71	8, 7
10	.	60.0	ហ	- ⊣	US-08-619-462-8	ထဲ
11	æ	60.09	5	~	68-	Sequence 36, Appl
12	τ.	60.09	5	3	80-60-	œ
13	m	60.09	S	4	-09-421-845-	w
14	Ψ.	0.09	5	4	US-09-298-924-35	1.,
15		60.09	5	9	5276016-4	No.
16	m	60.0	5	9	5276016-8	Patent No. 5276016
17	ε:	96	u i	(4)	5276016-10	Piten* No. 5276916
18	₹,	=: =:	G.	٠	5	Patent No. 5276016
16	m	60. n	·£)	4	2-698-	Engineers 40, Apply
20	с*	60.0	٤	-	2	,£
21	æ	60.0	9	r4	US-08-137-117D-18	۲,
2.2	m	(0.09	Ψ	C1	-08-4	Sequence 137, App
23	ρŃ	60.03	9	-4	-08-436-0	Sequence 3, Appli
24	~	60.09	_	2	US-08-702-673-4	Sequence 4, Appli
60. CV	6	0.09		64	-08-803	~
26	m	60.09	9	7	US-08-310-912A-13H	Sequence 138, App
27	æ	60.0	S.	4	US-09-113-921-44	Sequence 44, Appl

77	0.09	o	.	US-08-476-134A-33	
e)	60.0	ų)	÷	US-09-301-085-138	Sequence 1:8, App
~	0.09	ت		02-09-302-3050-8	Sequence P. Appli
~	0.09	9	ur,	PCT-US93-01669-30	Sequence 30. Appl
,~ ,	60.0	9	ın,	PCT-US93-01669-31	Sequence 11. Appl
m	60.0	٠2٠	ď	PCT-US95-04589-148	Sequence 138. App
æ	0.09	VÇ.	s.	5190920-6	Patest No. 5195920
er.	0.09	w	÷	5200320 15	Patent No. 5200320
m	60.0	9	9	5506208-6	Patent No. 5506208
m	0.09	7	-	US-08-201-046A-13	Sequence 14. Appl
m	0.09	,	-	US-08-076-092-34	Sequence 33, Appl
m	0.09	ŗ-	~	US-08-318-970R-16	Sequence 16. Appl
~.	60.0	۲÷		US-07-942-245-501	Sequence 501. App
(m)	0.09	7	2	US-08-483-636-18	Sequence 18. Appl
~	0.09	r ·	\sim	US-08-483-632-18	Sequence 18 Aprl
(ar.	0.00	ř-	-7	05-08-730-486-34	Sequence 33. Appl
ر~`	0.09	r-	C4	US-08-645-193B-31	Sequence 41. Appl
æ	60.0	7	٣,	US-08-335-865J-15	Sequence 15, Appl

ALIGNMENTS

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Sequence 247, Application US/09177249

Sequence 247, Application US/09177249

Patent No. 6229064

GENERAL INFORMATION:
AFFLICAUT. Fischer. Commonity AFFLICAUT. Rivosue, Tomobiro AFFLICAUT. Rivosue, Tomobiro AFFLICAUT. Margossian, Linda AFFLICAUT. The Ference of Control Seed and Fruit AFFLICAUT. The Ference of Control Seed and Fruit TILE OF INVENTION: Nucleic Acids That Control Seed and Fruit TILE OF INVENTION: Development in Plants
FILE REFERENCE: 023070-0861200S

CUPPENT FILING DATE: 1999-10-22

EARLIER APPLICATION NUMBER: US 09/071,838

EARLIER APPLICATION NUMBER: US 09/071,838

EARLIER PLING DATE: 1998-05-01

NUMBER OF SEO ID NOS: 324

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRESENT 2

URS-08-385-443-5

Sequence 5. Application MS/08385443

Faterin No. 5534455

GENERAL INFORMATION:
AFFLICANT: RUFF, MICHAEL R.
AFFLICANT: RUFF, MICHAEL R.
ITLE OF INVENTION: FAIN AYMORGANES

NUMBER OF SEQUENCES: 6

CORRESPONDENCE B. GOLDHERG

SIRRET: 261 MADISON AVENUE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 3; DB 4; Length 4:
; Pred. No. 20+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 60.0%; Score 3; DR 4 Best Local Similarity 100.0%; Pred. No. 2e+0 Matches 3; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ORGANISM: Arabidopsis sp. US-09-177-249-247
RESULT 1
US-09-177-249-247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 SYT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 SYT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGIH: 4
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REGISTRATION NUMBER: 34,409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE:
                                                                                                                                IYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Taddei-Peters, W. C.
APPLICANT: Butler, Sandra M.
TITLE OF INVENTION: Immunoreactive Peptides of Apo(a)
NUMBER OF SEQUENCES, 48
COKRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                        REFERENCE TOON NUMBER: 24,408
REFERENCE TOOCHER NUMBER: ADVANCED PEPTIDE: PS
TELECHMONICATION INFORMATION:
TELEPHONE: 212-818-9479
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACIERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PatentIn Release #1.0, Version #1.25
                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  60.0%; Score 3; DB 1
ilarity 100.0%; Pred. No. 2e+
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 01-UUN-1995
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/266 'CT
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER, US 08/172,461
FILING DATE: 21-DEC-1993
ATTOENEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 08/067.523 FILING DATE: 23-MAY-1993 AITCHEYAGENT INFORMATION:
                                                                                                                                                                                       US/OR/385,443
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DGS/MS-DGS
                                                                                              E: Floppy disk
IBM PC compatible
SYSIEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 36, Application US/U8456840
Patent No. 5597908
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Akzo No. 5597908el
STREET: 1330 Piccard Drive
CITY: Rockville
SIATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SS: not relevant not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 27-JUN-1994 APPLICATION NUMBER. US 08
                                                                                                                                                                                     APPLICATION NUMBER: US/ORFILING DATE: 08-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gormley, Mary E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21P: 20850
COMPUTER READABLE FURM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: not relevan
                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
CITY: NEW YORK
STATE: NEW YORK
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                                                                                                                 COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-456-840-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 SYT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-385-443-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 SYT 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
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9170
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Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 3: Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60.0%; Score 3; DB 1; Length 5; 100.0%; Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: HS//W/170,360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5. Rothwell, Figg Ernst & Kurz
Suite 701-E, 555 Thirteenth St., N.W
                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Tseng, Albert P. S.
APPLICANT: Inglis, Adam
TITLE OF INVENTION: PLA2 INHIBITORY COMPOUNDS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION UMREP: 41 PT/AU72/00333
FILING DATE: 06-JUL-1992
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
RILING DATE: 04-JUL-1991
ATTORNEY/AGENT IRPERMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-MAR-1994 N. 514 N. 717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Ernst, Barbara G. PESISTRATION NUMBER: 30.377
PESISTRATION NUMBER: 3871-104A
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11, Application US/08170360
Patent No. 5656602
TELECOMMUNICATION INFORMATION:
TELEPHONE: 301-258-5200
TELEFAX: 301-977-0847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEÓ ID NO: 11:
SEQUENCE CHAPACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (202)783-6040
(202)783-6040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; FRAGMENT TYPE; N-terminal US-08-170-360-11
                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 amino acids
                                                          INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                 , MOLECULE TYPE: protein US-08-456-840-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 20004
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Washington
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Best Local Similarity
                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                          RESULT 4
US-08-170-360-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                           1 LES 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET:
```

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APPLICANT: RATHJEN Deborah
APPLICANT: WIOMEN, Fred
TITLE OF INVENTION: Pertine T and Pelated Peptides in the
TITLE OF INVENTION: Treatment of Inflammation, Including Multiple Scienosis
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        60.0%: Score 3: DB 1: Length 5: 100.0%: Fred. No. 2e+05; tive 0: Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                           COMPUTER: IRM PC compatible
OFERATING SYSTEM: FC FOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION BIMPER: TS F0 730, 979
FILING DATE: 29-MAR-1993
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptides for Treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION DATA:

ALFICATION DATA:

ALFICATION DATA:

ALFICATION DATA:

PILING DATE: 99 MAR-1993

PROR APPLICATION DATA:

AFELDATI WINNEFF IN: "197,674

FILING DATE: 99 DEC-1992

PROR APPLICATION DATA:

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APPLICATION DATA:
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ADDRESSEE: BANNER & ALLEGRETTI, LTD
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PLING DATE: 14-MAY-1992
PRIOR APPLICATION DATE: 12-07,950,932
                 DOOB, Penelope R
MacFADDAEN, Douglas K
PHIPPS, David J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 6, Application US/08481840A; Patent No. 5763406
                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PREPERSTOCKET NUMBER: 91
TELECOMMUNICATION INFORMATION: PELEPHONE: 312-715-1000
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                                                                                                                                                                                                           10 S Warker Drive
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ATTORNEY, ASENT INFORMATION:
NAME: MCDOODER! John J
 CARLEN, Peter L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: REDERSEN, OVE
APPLICANT: MACFADDEN, D.
TITLE OF INVENTION: Bearti
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Matches 3; Conservative
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MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER:
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                                   APPLICANT:
APPLICANT:
                     APPLICANT:
   APPLICANT
                                                                                                                                                                                                             STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CEEATINY SYSTEM: PT. (**) MS. (**) MS. (**) SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                           APPLICANT: TSend, Albert P. S.
APPLICANT: Indils, Adam
ITILE OF INVENTION: PLAZ INHIBITORY COMP.GNUS:
CORRESPONDENCES: 17
CORRESPONDENCE ADDRESS:
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100.0%; Pred. No. 2e+0';
tive 0; Mismatches
O: Mismatches
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APPLICATION NUMBER: AU PCT/AUG2/On333
FILING DATE: 05-JUL-1992
PPLOR APPLICATION DATA:
APPLICATION NUMBER: AU PK 7058
FILING DATE: 04-JUL-1991
ATTORNEY, AGFNT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1871-104A
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CLASSIRICATION: 514
RIOR APPLICATION: 514
                                                                                                                                                                     Sequence 14, Application Hs/OR17OANU
Patent No. 5656602
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Patent No. 5756449
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APPLICANT: ANDERSEN, Anders J
APPLICANT: ASION, Roger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Ernst, Barbara G.
REGISTRATION VUMBER 30,377
REFERENCE/TOOPKET NUMBER 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: (202) 793-F040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-terminal
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 Conservative
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Best Local Similarity
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                                     3 SYT 5
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                US-08-266-407A-36
                                                                                                                                                                                           TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Taddei-Peters, W. C.
APPLICANT: Buller, Sandra M.
TITLE OF INVENTION: Immunoieactive Peptides of Apo(a)
NUMBER OF SEQUENCES: 48
AUGHESSEE: Akzo No. 5786156e1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                    ZIP: 606.05
ZIP: 606.06
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk, 3.50 inch
COMPUTER: IBM PC compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/481.845A
FILING DATE: 07-JUN-1995
CLASSIFICATION: 514
        NUMBER OF SEQUENCES: 7
CORRESSONDENCE ADDRESS:
ADDRESSEE: Banner & Allegretti, ild.
STREET: 10 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/265,407A FILING PATE: 27-TMN-1994 CLASSIFICATION: 435 PRIOR APPLICATION DATA: APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                           1 1 2 34,628
PFR: 93,741-A
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FILING DATE: 21-DEC-1993
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SYSTEM: PC-DOS/MS-DOS
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atent No. 5786156
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/107,777
FILING DATE: 24-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 93
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-9110
TELEFAX: 617-345-9111
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INFORMATION FOR SEQ ID NO: 6:
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                                                                                                                                                                                                                                                                                                                                                                                                                   John P
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: Linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC (OPERATING SYSTEM:
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                                                                               CITY: Chicago
STATE: Illinois
COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MICHAELIS, JURGEN
APPLICANT: SLEIGH, MRRILYN J.
TITLE OF INVENTION: TREATMENT OR PREVENTION OF CPOHN'S
TITLE OF INVENTION: ISEASE AND,/OR !!CERATIVE COLITIS
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/403 718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
2IP: 22201-4714
COMPUTER READABLE FORM:
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
NAME: Gormley, Mary E.
REGISTRATION NUMBEP: 14,409
TELECOMUNICATION INFORMATION:
TELEPHONE: 301-258-5200
TELEFAX: 301-977-0847
INFORMATION FOR SEQ 1D NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/08403718
Patent No. 5795858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: MITCHARD, LEONARD C. ERSISTRATION NUMBER: 29,009 REFERENCE/DOCKET NUMBER: 47-TELECOMMUNICATION INFORMATION: TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (704) 816-4100 INFORMATION FOR SEQ ID NO: 78:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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GY: linear
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Gaps
                           Sequence 8, Application US/08619462
Patent No. 5798335
GENERAL INFORMATION:
APPLICANT: MICHAELIS, JURGEN
APPLICANT: Trigg Timothy bliggt
TITLE OF INVENTION: METHOD FOR THE TREATMENT HE PREVENTION OF
TITLE OF INVENTION: BCTEMA/PERMATITIS
NUMBER OF SEQUENCES: 8
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APPLICANT: Taddei-Feters, W. C.
APPLICANT: Butler, Sandra M.
TITLE OF INVENTION: Imminoreactive Peptides of Apo(a)
NUMBER OF SEQUENCES: 48
COPPESPONDENCE ALIPESS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTAM: PC-PASAMS-LOS
SOFTWARE: Patentin Release #1.0, Version #130
CURRENT APPLICATION DATA:
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COMPUTER: IBM PC Compatible
OPERATIVA SYSTEM: PC-EGS/MS-TGS
SUFIWARE: Patentin Release #1.0, Version #1.25
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mer: US;08;619,462
07-MAY-1996
N: 514
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STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Floppy disk
IBM PC compatible
SYSTEM: PC-DGS/MS-LGS
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Patent No. 5874544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: MITCHARD, LEDNARD C.
REGISTRATION NUMBER: 29,004
REPERBORE/SOCKET NIMBER: 47-84
TELECOMMUNICATION INFORMATION:
TELEFANE (704) 816-4000
INFORMATION FUR SEQ ID MG: 8:
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21F: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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FILING DATE: 07-MAY
CLASSIFICATION: 514
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US-08-619-462-8
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TITLE OF INVENTION: Perolide T and Related Peptides in the Treatment of TITLE OF INVENTION: Inflammation, Including Multiple sclerosis NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
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jin i*; Fred No 20+05;
tive 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy dish
COMPUSE: THM EC COMPATIBLE
COMPUSE: THM EC COMPATIBLE
COMPANIES: PATCHER FELICS/MS FOS
SOFTWARE: PATCHER Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
AFFLICATION NUMBER: US/US/082/837A
FILING DATE: 21.3UH 1938
TYPER CATION: 514
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jord*; Fred a.
vo 0; Mismatches
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FILING DATE: 29-MAR-1993
PRIOK APPLICATION DATA:
APPLICATION NUMBER: US 277987, 674
                                                                                                                   FILING DATE: 21-DEC-1993
ATTORNEL/AGENT INFORMATION:
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APPLICATION TUMBER: US 08/202,829
FILING DATE: 29-MAR-1993
                                                                      PRIOR APPLICATION DATA:
APPLICATION PYMEE: 12,12,12,126, 407
FILING DATE: 27-7UN-1994
APPLICATION PYMBEE: 08-02,172,461
                 US, rop, /892, 544
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10 S. Wacker Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ANDERSEN, Anders J
                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
IELEPHONE: 301-258-5200
TELEFAX: 301-977-0847
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHIPPS, David J
RATHJEN, Deborah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DOOB, Penelope R
MacFADDAEN, Dougl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ASTON, Roger
CARLEN, Peter L
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CURRENT APPLICATION DATA: APPLICATION NUMBER: US FILING DATE:
                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO.
                                                                                                                                                                                                                                                                                                                      5 amino acids
                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein
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                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                      CLASSIFICATION:
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US-09-082-837A-8
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1 LES 3
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APPLICANT:
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APPLICANT:
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FILING DATE:

us-09-647-749a-1.oliszlm.rai

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Sequence 35, Application HS,00208924 Patent No. 6391595 GENERAL INFORMATION:
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Best Local Similarity
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;Patent No. 5276016
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Pred. No. Ze+∪
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                          US 07/915.118
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100.0%; Pre
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Patent No. 6265374
                                                                           FILING DATE: 14-MAY-1992
FRICK APPLICATION DATA:
APPLICATION NUMBER: US 07/858,
FILING BATE: 27-MAR-1992
ATTORNEY/AVENT INFORMATION:
NAME: DEERKOAFF, W. Definits
REGISTRATION NUMBER: 27193
FEREPROTE/FONCKET NUMBER: 27193
TELECHONE: 312-715-100
                                                                   DK 645/92
                                                                                                                                                                                                                                                                                                   not relevant
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                                                                                                                                                                                                                                        INFOEMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS:
                          APPLICATION NUMBER: US 07 FILING DATE: 17-JUL-1992
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                                                                                                                                                                                                                                                                                                                                                                                                         3; Conservative
                                                                                                                                                                                                                                                                         5 amino acids
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MOLEGULE TYPE: peptide
                                                    PRIOR APPLICATION DATA: APPLICATION NUMBER: 1
             PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: not
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Rathjen
Widmer
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US-09-298-924-35
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US-09-421-845-8
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APPLICANT:
APPLICANT:
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APPLICANT:
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KOBAYASHI, KAZUO
KOMEDA, TOShihiro
FULLE OF INVENTION: NOVEL TRANSPERASE AND AMYLASE, PROCESS
FOR THE SAME
FOR THE SAME
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                                                                                                                                                                                                                                                                                                             COMPUTE TOTAL
COMPUTER FEADMALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: TBM PC compatible
COMPUTER: PSYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
ARLHCALTAN NUMBER: REPORTING PATENTING DATE: 25-Apr-1999
CLASSIFICATION: CORROWN?
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ADDRESSEE: Foley & Lardner
STREET: 3000 K Street, N.W., Suite 500
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REFERENCE/DOCKET NUMBER: 49441/110
TELECOMMUNICATION INFORMATION:
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FILING DATE: UDKNOWN>
APPLICATION NUMBER: 07 7-120673
FILING DATE: 21-AFR-1995
APPLICATION NUMBER: JP 6-311185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: JP 6-290394
FILING DATE: 31-0CT-1994
APPLICATION NUMBER: JP 6-194223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 21-NOV-1994
APPLICATION NUMBER: JP 6-286917
FILING DATE: 21-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 35:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IELEPHONE: (202)672-5300
                                                            IWAMATSU, Akihiro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (202)672-5399
TELEX: 904136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ALTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stephen A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGIH: 5 amino acids
                    MIURA, Yutaka
KETTOKU, Masako
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMALIGN FOR SEG ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SIRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICANT: KATO, Masaru
MIURA, Yutaka
                                                                                                                                                                                        NUMBER OF SEQUENCES: 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Conservative
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## PAPLICANT: PERT, CANDACE B.;RUFF, WILLIAM F.
| FARRAW WILLIAM I. |
| ITLE OF INVENTION: SMALL PEPTIDES WHICH INHIB::
| BINDING TO T-4 RECEPPORS AND ACT AS IMMUNGENS |
| WINDER OF SEQUENCES: 19 |
| CURRENT APPLICATION DATA: |
| APPLICATION DATA: |
| APPLICATION DATA: |
| APPLICATION DATE: 14,507 |
| FILING DATE: 17-FEB-1999 |
| FILING DATE: 17-FEB-1999 |
| APPLICATION NUMBER: 314,507 |
| FILING DATE: 17-FEB-1999 |
| FILING DATE: 10-FEB-1999 |
| FILING DATE: 10-FEB-1999 |
| FILING DATE: 10-FEB-1999 |
| FILING DATE: 20-JUN-1996 |
| FILING DATE: 30-JUN-1996 |
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2792, Ap
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11, Appl
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Appl
Appl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Apsence 3, Ap
                                                                                                                                                                    (without alignments)
9.543 Million cell updates/sec
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                                                                                                                                             : Search time 21 5833 Seconds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and is derived by analysis of the total score distribution.
GenCore version 5.1.4_F%_4578 (2) 1992 - 2003 | Ourplan Etd.
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2. .gmz_1 [**d*fa, l. iaa, fh_cwmb.pcp.*
3. .m., p. ivalata, i. aa, ba_cwmp.p. i. a
4. .m., p. ivalata, i. aa, ba_cwmp.p. i. a
5. .m., prodata, laa, ba_cwmp.pcp.
6. .m., prodata, l. iaa, backtiine [Feb.
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                                                                                                                                                                                                                                                                                                                                                                                                                      262574 seqs, 29422922 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Post-processing: Listing first 45 summaries
                                                                                                 OM protein - protein search, using sw model
                                                                                                                                             April 29, 2003, 98:50:15
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                         Cupyrisht (c) 1992 -
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Match Length DB
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                                                                                                                                                                                                                                                                           Perfect score:
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                                                                                                                                                                                                                                                                                                     sednence:
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Sequence 18 Appl
Sequence 457, Appl
Sequence 457, Appl
Sequence 1037, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 12, Appl
                                            41, Appl
52, Appl
4, Appli
5188642
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APPÌ
                                                                                                                                           App
                                                                                                    Patent No. 5310667
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Blaschuk, Orest W.
APPLICANT: Gour, Barbara J.
TILLE OF INVENTION. COMPOUNS AND METHODS FOR MEDIATED AND MEDIATED FOR MEDIATED FUNCTIONS
TILLE OF INVENTION. CARBERTY MEDIATED FUNCTIONS
FILE FERFERPY: J. 0gt 47021
"UPFERT FILIAGONE: 1998-11-06
TUPFERT FILIAGONE: 1998-11-06
SPPTWARR: PALENTIN Ver. 2.0
SPPTWARR: PALENTIN Ver. 2.0
                                                                                                                                           nes actambés
                              Sequence 41,
                                                         Sequence 52,
                                                                                                               Sequence 31
    Sequence 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Pepresentative cyclic modulatina agent based on OTHER INFORMATION: Protocadherin cell adhesion recognition sequence
                                                                        Sequence 4.
                                                                                                                              Sequence 5
                                                                                      Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: The Scripps Research Institute
TITLE OF INVENTORS OF TWOF-SPECIFIC
TITLE OF INVENTORS OF OFFICE OF TWOF-SPECIFIC
TITLE OF INVENTORS OF OFFICE OFFICE
COPPRESSORDERS ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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US 33 243 374 28
US 99 623 6188 6
US 08 444 000 41
US 08 445 000 41
US 08 55 605 52
PCT US95 10793 4
510657 19
US 08 137 514A 31
US 08 137 514A 18
US 08 137 514B 18
US 09 624 17 310A 18
US 09 625 10793 5
US 09 555 10793 5
US 09 555 10793 5
US 09 556 605 51
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STREET: Tokes Morth Torieg Pines Boal, TRC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 4; DB 4
Pred. No. 2e+
0; Mismatches
                                                                                                                                                                                                                                                                                    AL LUNMENTS
                                                                                                                                                                                                                                                                                                                                         US-09-187-859-2791
; Sequence 2791, Application US/09187859A
; Putent No. 6358920
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 2
PCT-0292-16415-34
PCT-03-16415-34
PCT-03-16415-34 PCT-03-16415
PCENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       57.18;
100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READARLE FORM: MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: La Jolla
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                    .; GENERAL INFORMATION:
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| EYFT 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGIH: 9
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Sequence 41,

US-08-297-731-4 US-08-290-448A-41 US-08-290-448A-41 US-08-290-448A-41 US-08-578-694A-41 US-08-578-573-4 US-08-893-6548-13

PCT-US95-13794-20 PCT-US95-13794-71

Sequence Sequence Sequences Sequence Sequence Sequence Sequence Sequence

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Sdeb
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APPLICANT: Blaschuk, Orest W.
APPLICANT: Blaschuk, Orest W.
TITLE OF INVENTION - COMPOUNDS AND METHODS FOR M HITATING NONCLASSICAL
FILLE OF INVENTION: CAMPBINIMEDIATED FUNCTIONS
FILE REFERENCE: 100086.407c1
CUPRENT APPLICATION NUMBER - US/03/187.859A
CUPRENT FILING DATE: 1998-11-06
SUPPRENT FILING DATE: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Expressitative cyclic mechanisms agent based on OTHER INFORMATION: Protocadherin cell adhesion recognition sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P. Attento
                SOFTWARE: Patentin Release #1.0, Version #1..5 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query March 57,14, South 4, TB 5, Best Local Similarity 100,0%; Pred, No. 2e+05; Matches 4; Conservative 0; Mismatches
                                                           MBER: PCT/US95/16415
13-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2792, Application US/09187859A Patent No. 6358920
                                                                                                                                                         APPLICATION NUMBER: US 08,7355,558
                                                                                                                                                                                                                                                                       REFERENCE/INCRET NUMBER 433 1PC TELECOMMUNICATION INFORMATION:
OPERATING STSIEM. FC-DOS/MS-LOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vincent, Jean-Pierre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 3, Appli, atlum US,08416007; Patent No. 5693679
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Beaudet, Alain
                                                                                                                                                                                                                                                                                                             TELEPHONE: (619) 554-2937
TELEFAX: (619) 554-6312
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                 33,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artiticial Sequence
                                                                                                                                                                         FILING DATE: 14-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                            NAME: LOGAN, April C. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide PCT-US95-16415-34
                                                                                                           CLASSIFICATION:
PRICE APPLICATION DATA:
                                                              APPLICATION NUMBER:
FILING DATE: 13-DEC
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APPLICANT: Vincent
APPLICANT: Gaudria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CS-08-187-859-2792
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LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              * EYFT 6
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                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30
CHPRENT APPLICATION DATA:
AFFLITATION WINNER: NS/09/416 007
FILING DATE: 04-APR-1995
CLASSIFICATION: 530
ATTORNEY/ASCENT INFORMATION:
NAME: Clark, Paul T
REGISTRATION, NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1 30
CURRENT APPLICATION DATA:
APPLICATION NUMBEP: IS.08/475,751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Petent No. 582472

CENERAL INFORMATION US/08475751

PETENT NO. 582472

CENERAL INFORMATION

APPLICANT: Gaudriault, Georges

APPLICANT: Gaudriault, Georges

APPLICANT: Baundet, Alain

TITLE OF INVENTION

FUNDRESPORTS

CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

STREET: 585 COMMERCIAL Street
TITLE OF INVENTION. FLORESTENT SCHAIGSTATIN NUMBER OF SEQUENCES: 5 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
42.9%: Score 3: DB 1;
Best Local Similarity 100.0%: Pred. No. 2e+(
Matches 3: Conservative 0: Mismatches
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ne942 me9901
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CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/416.007
FILING DATE: 04-APR-1995
                                                         Elsh & Richardson P.C.
225 Franklin Street
                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                   FEFERENCE, TOOKKET NUMBER 1867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           not relevant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                        : 617/542-5070
617/542-8905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 200154
INFOFMALION FOR SEQ 1D NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIF: 02109-1024
                                                                                                                                  Y: USA
02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Boston
                                                                                                Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE:
                                                                                                                  STATE: MA
                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX:
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                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-416-007-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY:
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STRANDEDNESS: single

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DB 2,
_2e+05;
`-~ 0; Indels
                                                                                                                                                                                                                                                                                                                                                                              Length 4;
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APPLICANT: Bergmann, Carl
TITLE OF INVENTION: Nucleotide Sequences Coding An
TITLE OF INVENTION: Endopolygalacturonase incubitor
NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patenti Coop.m. 25
SOFTWARE: Patenti Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER 118 (9-144 644
                                                                                                                                                                                                                                                                                                                                                                         Query Match 42.9%; Score 3; DB 2;
Best Local Similarity 100.0%; Pred. No. 2e+05
Matches 3; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Sally A. Sullivan STREET: 5370 Manhattan Circle Sulte 201
                NAME: Clark, Paul T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 06942,004nn1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617,723.4123
TELEFAX: 617,723.48952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: IT PM 91A 000915
FILING DATE: 06-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/IT/00158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-FOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 19-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3. Application HS/00244646
Patent No. 5744692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGENT Internal Sullivan, Sally A. Sullivan, Sally A. Timmerr, 32,064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Cervone, Felice
APPLICANT: De Lorenzo, Giulia
APPLICANT: Salvi, Giovanni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: (303)499-8080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Salvi, Giovanni
APPLICANT: Albersheim, Peter
                                                                                                                                                                                                                                                                   not relevant
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INFORMATION FOR SEQ 1D NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 06-JUN-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 04-DEC-1992
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 4 amino acids
                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COPPESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Boulder
                                                                                                                                                                                                                                                            STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
                                                                                                                                                                                                                                              TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 6
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APPLICANT: Madeen, Made Thorup
IIILE OF INVENTION: An enzymatic process for
TILIE OF INVENTION, predacing a dustred prittin from an amin, terminally extend
NUMBER OF SEQUENCES: 17
                                                                                                                                                                     Query Match
42.9%: Score 3: DB 1: Length 5:
Rest Local Similarity 100.0%; Pred. No. 2e+05;
Matches 3: Conservative 6: Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42.0%; Score 3; DB 1; Length 5; 100.0%; Fred, No. 2e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER. IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEG for Windows Version 2.0
CURRENT APPLICATION DATA:
AFPLICATION NUMBER 195/797,737,757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Darby & Darby. PC
STREET: 805 Third Avenue - 27th Floor
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4.11,00596
                                                                                                                                                                                                                                                                                                                                            RESULT 7
US-08-737-757-3
Sequence 3. Application US/08737757
Patent No. 5783413
                                                                                            ORGANISM. Phaseclus vuigaris
STRAIN: Pinto
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TELECOMMUNICATION INFORMATION:
TELEFRONE: 212-527-7700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lauritzen, Conni
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMFUTER. IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEYZAGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                             Pedersen, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Obery Match
Bost Local Similarity 100.0
"has 3, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 5 amino acids amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: GOGOTIS, Adda C
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
TOPOLOGY: linear
            MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: peptide US-08-737-757-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10022-7513
                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                          ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USA
                                      HYPOTHETICAL: NAMANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
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    TOPOLOGY:
                                                                                                                  ; STRAIN;
US-08-244-646-3
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2 FTS 4
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                                                                                                                                                                                                                                                   5 FTS 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 FTS 7
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
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RESULT 10
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                                                                                                    APPLICANT: Jackyoon Shin, Insil Joung, Rather K. Vadlamudi APPLICANT: and Jack L. Strominger TITLE OF INVENTION: p62 POLYPEPTIDES, RELATED POLYPEPTIDES IITLE OF INVENTION: AND USES THEREFOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            o, Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42.9%; Score 3: DB 2: Length 5; 100.0%; Pred. No. 2e+05. ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                       ZIP: 02109-1875

ZIP: 02109-1875

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
COMPUTER: IBM FC compatible
COMPUTER: STEEL PC-TORS/MS-DOS
SOFTWARE: Patentle Release #1.0. Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEFFENCY INVEST NUMBER OF OF OR TELECOMMUNICATION INVESTMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-5941
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH : 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APELICATION HYMBER (13,008,574,959A FILLING DATE: 19-DEC-95 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                   NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEB: ALAIVE & COCKFIELD
SIREET: 60 State Street, Suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: NIXON & VANDERHYE P.C. STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/08432016
Patent No. 5968768
GENERAL INFORMATION:
APPLICANT: HAYNES, BARTON F.
APPLICANT: ARUPFO, ALEDANDRO
APPLICANT: PATEL, DHAVALKUMAR
APPLICANT: BOWEN, MICHAEL A.
APPLICANT: MARQUARDT, HANS
                                        Sequence il, Application OS/vo574354A
Patent No. 5962224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD6 LIGAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Mandragouras, Amy E. REGISTPATION NUMBER: 36,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 100.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 22201-4714
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                    STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: ON NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: ARLINGTON STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         USA
                                                                                                                                                                                                                                                                                            Bustor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-574-959A-11
RESULT 8
US-08-574-959A-11
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US-08-432-016-7
                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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Query Match 42.9%: Score 4: DB 2: Length 5: Best Local Similarity 100.0%; Prod. No. 2e+05: Matches 3: Conservative 0: Mismatches 0: Indels
                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CUPPENT APPLICATION DATA:
AFFLICATION NAMES: NS/44/2/016
FILING DATE: 01-May 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 05-NOV 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NAMER: 18,708/684,594
FILING DATE: 18-JUL-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: HS 08/432,016
FILING DATE: 01-MAY-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: HS 08/333,350
FILING DATE: 02-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: NIXON & VANDERHYE P.C. 1100 NORTH GUERE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 1579-95
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 816-4000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IRM PC compatible OPERATING STSTEM: PC:DOS/MS-DOS
MEDIUM IYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 7, Application US/08684594
, Fatent No. 5998172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: CD6 LIGAND NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PATEL, PHAVALKHMAR
                                                                                                                                                                                                                                                                                                                                                                                    32,955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: HAYNES, BAKTON F.
APPLICANT: ARGFFO, ALEJANDRO
APPLICANT: PATEL, DHAVALK'HMAR
APPLICANT: BOMEN, MICHAEL A.
APPLICANT: MARQUARDT, HANS
                                                                                                                                                                                                                                                                                                        FILING DATE: 02-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGISTRATION NUMBER: 32.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (703) 816-4100
INFORMATION FOR SEQ 1D NO: 7
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; MOLECULE TYPE: protein 38-08-432-016-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LITY: ARLINGTON STATE: VIPOT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 IKE 3
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2, 345
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TITLE OF INVENTION: Interactions and Methods of Use Therefor
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS.
ADDRESSER: HAMILLOG, Brock Smith & Prynoids p.c.
STREET: Two Militia Drive.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            3. Tride() 5.
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 42.9%; Score 3; DB 2; Longth 5; Best Lonal Similarity 100 0%; Pred No 20+06. Matches 3, Conserrating 0, Mismatches 1, Tobalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-LT-S/MS-LT-S
SOFTMARE: Patentin Release #1.0, Version #3:+0
CUPPENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: /label+ modified as OTHER INFORMATION: /note → "Ac - Leucine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Brook, David E.
REGISTRATION NUMBER: 12,592
REFERENCE/DOCKET NUMBER: 1KS95-12
TELECOMMUNICATION INFORMATION:
                                                                                                                           REFERENCE/DOWNET NUMBER: 1579-112
TELECOMMUNICATION INFORMATION:
TELEPHONF: (703) 816-4000
PRIOR APPLICATION LATA:
APPLICATION NUMBER: IS OR/141,903
FILING DATE: 02-NOV-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US, 08, 582, 740
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/ Sequence 42, Application US/08582740
/ Patent No. 6037324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Schwender, Charles F. APPLICANT: Shroff, Hitesh N.
                                                                                                       REGISTRATION NUMBER: 32,955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (617) 861-6240
TELEFAX: (617) 861-9540
INFORMATION FOR SEQ ID NO: 42:
                                                                                                                                                                                            TELEFAX: (703) #16-4100
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JAN-1996
1: 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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.. Massachusetts
RY: USA
                                                                                                                                                                                                                                                                5 amino acids
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amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                         WILSON, MAPY J
                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02173
COMPUTER READARLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AEPLICATION NUMBER.
FILING DATE: 04-JAN
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                  amino acid
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                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                     US-08-584-594-7
                                                                                                                                                                                                                                                           LENGTH
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                                                                                         NAME .
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TITLE OF INVENTION: PGLYMORPHORUGLEAR LEUKOCYTE ELASTASE ACTIVITY
NUMBER OF SECUENCES: 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42.9%; Score 3; DB 3; Length 5; 100.0%; Pred, No. 2e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                  Query Match 42.9%; Score 3; DR 3; Length 5; Best Local Similarity 100.0%; Pred, No. 2e+05; Matches 3; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: YEM FC Compatible
OPERATING SYSTEM: PC-ToSAMS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 71
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: RE. CHEFSTINE E. CARTY
STREET: 126 E. LINDGLM AVENUE., P.O. BOX 2000
                                         LOCATION: 5
COTHER INFORMATION: /labei-modified aa
CTHER INFORMATION: /noter "Leucine : NH2"
US-08-582-740-42
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APPLICATION NUMBER: #5,08,469,141A
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                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20, Application US/08469141A
Patent No. 6124107
SENEFAL INFORMATION:
AFFLICANT: MTMFORE, FICHARD A.
APPLICANT: DAVIES, D.T. PHILIP
APPLICANT: DAHLGREN, MARY E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
ATHORNEY/AGENT INFORMATION:
NAME: CARTY, CHRISTINE E.
REGISTRATION NUMBER: 35,099
EEFEENIE/FOCKET NUMBER: 37467
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: DAHLGREN, MARY E. APPLICANT: BGGER, JOSHGA S. APPLICANT: HUMES, JOHN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (908) -594-6734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION F04-4720
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                      NAME/FEY: Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-469-141A-20
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                                                                                                                                                                                                                                            5 FTS 7
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FEATURE:
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Gaps
                                                                                                APPLICANT: MUMPORD, RICHARD A.
APPLICANT: DAVIES, D.T. PHILIP
APPLICANT: DAHLGREN, MARY E.
APPLICANT: DAHLGREN, MARY E.
APPLICANT: BOGEN, JOHN L.
APPLICANT: HUMES, JOHN L.
TITLE OF INVENTION: POLYMORPHONUCLEAR LEUKOCYTE ELASTASE ACTIVITY
NUMBER OF SEQUENCES: T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Letner, Righard A.

TITLE OF INVENTION: SYNTHETIC HUMAN NEUTRALLLING MONOCLONAL

TITLE OF INVENTION: ANTIBODIES TO HUMAN IMMON-DEFICIENCY VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: The Scripps Research Institute, office of ADDRESSEE: Patent Counsel STREET: 10550 No. 6261558th Torrey Pines Read, TPC 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 42.9%; Score 3; DB 3; length 5; Best Local Similarity 100.0%; Pred. No. 2e+55. Matches 3; Conservative 0; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 126 E. LINCOLN AVENUE., P.O. BOA ACTO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRENT AFFILIATION ADDRESS

COASSIFICATION: 40 C-JUN-1995

CLASSIFICATION: 40 C-JUN-1995

ATTORNEY AGENT INFORMATION:
NAME: CARTY, CHRISTINE E.
REGISTRATION NÜMBER: 36,099

REFERENCE/POCKET NÜMBER: 174611B

TELEDOMMUNICATION INFORMATION:
TELEPAN: (908)-594-6734

TELEFAX: (908)-594-4720

INFORMATION FOR SED, 10 NO: 71:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                             DR. CHRISTINE E. CARTY
                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
OVERATING SYSTEM: PC-DOS/MS-DOS
                                     Sequence 71, Application US/08469141A Patent No. 6124107 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24, Application US/08591632
Patent No. 6261558
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Dennis R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                          NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                                                                                                                                                                                                                  CITY: RAHWAY
STATE: NEW JERSEY
COUNTRY: USA
ZIP: 07065-0907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Burton,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Barbas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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APPLICANT: Barbas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       La Jolla
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                                                                                                                                                                                                                                                                                               ADDRESSEE:
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                     US-08-469-141A-71
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CITY: R.
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hamilton, Brook, Smith & Reynolds, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1 0, Version #1.30 CURPENT APPLICATION DATA: APPLICATION NUMBER: 02,09,109,879 FILING DATE: 02-JUL-1998 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Hitch 42.9%; Score 3; DB 4; L
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 3; Conservative 0; Mismatches 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRING DATE: 19-0CT-1994
PRIOR APPLICATION DATA:
APPLICATION HUMBER: 15-0CF-1994
FILING DATE: 19-SEP-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 15-0CF-1994
PRIOR APPLICATION NUMBER: 15-0CF-1994
PRIOR APPLICATION NUMBER: 15-0CF-1994
PRIOR APPLICATION NUMBER: 15-0CF-1994
APPLICATION
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                                                                                                                                                                                                                                                                                                                                                                                                  AFILS A.E. 19-0CT-195-
PRING DATE: 19-0CT-195-
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/11907
                                                                                                                                                                                                                                                                                                                                                  CORRENT APPLICATION DATA:
AFFLITATION NUMBER - NEWS 1591,(32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IRM PC compatible
OPERATING SYSTEM: PC-DOS/MS-D/S
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-109-879-42; Sequence 42; Sequence 42; Partent No. 6274556; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   34,163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: TS
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (619) 784-2937
TELEFAX: (619) 784-9399
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Two Militia Drive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 19-0CT-1993
ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fitting, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGIH: 5 amino acids
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CITY: Lexington
STATE: Massachusetts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 8'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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USA
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                                                             COUNTRY:
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PRICK APPLICATION DATA:
APPLICATION NUMBER: 1977/1947/104291
FILING DATE: 03-378-197
FILING DATE: 04-388-196
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 25,592
REGISTRATION NUMBER: 25,592
REGISTRATION NUMBER: 25,592
REGISTRATION NUMBER: 26,400
INFORMATION FOR SEG ID NO: 42:
FELEPAN: (781; 861-954)
INFORMATION FOR SEG ID NO: 42:
FELEPAN: 10 861-6540
INFORMATION FOR SEG ID NO: 42:
FELEPAN: 10 861-6540
INFORMATION FOR SEG ID NO: 42:
FELEPAN: 10 861-6540
INFORMATION FOR SEG ID NO: 42:
FELEPAN: 10 861-6540
INFORMATION FOR SEG ID NO: 42:
FENTURE: INFORMATION: /label = modified as office INFORMATION: /label
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Search completed: April 29, 2003, 08:55:04 Job time : 23.5833 secs

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GenCore version 5.1.4_pt.4578

Copyright (c) 1993 - 2003 Compagne Ltd.

OM protein - protein search, using sw model

Fun cor April 20, 2003, 00.35.29 ; Search - --- 27 4167 Seconds

(without alignments)

52.608 Million cell updates/sec

Title:

Perfect score: 7

Sequence: 1 IKEYFIS 7

Scoring table: OLIGO
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Sequence: 1 IMEMPTS 7

Scoring table: OLIGO Gapop 50.0 , Gapext 60.0 Gapop 50.0 , Gapext 60.0
Searched: 671580 seqs, 206047115 residues
Word size : 0

Total number of hits satisfying chosen parameters:
Minimum DB seq length: 0

Maximum DB seq length: 10

Post-processing: Listing first 45 summaries
Database : SPTREMBL_21:*
1: Sp_archea:*

SPTREMBL_21:*	1: sp_archea:*	 sp_bacteria.* 	3: sp_fungi:*	4: sp_human:*	5: sp_invertebrate:*	€: ⊱p_mammal:*	7: sp_mhc:*	8: sp_ordanelle:*	9: sp_phage:*	10: Sp_plant:*	11: sp_rodent:*	12: sp_virus:*	<pre>13: sp_vertebrate:*</pre>	14: sp_unclassified:*	15: sp_rvirus:*	17: sp archeap:*
base :																

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Qun6m5 toxoplasma	Q934s1 thermus the	Q9np07 homo sapien	திவந்தி நின்ற தவநித்	P72081 nocardia la	19s6d5 escherichia	Q9r7t2 escherichia	ns sultipad jerihij	Q56759 xanthobacte	P77556 escherichia	Q8rji0 streptomyce	P87225 saccharomyc	Q9h4d3 homo sapien	Q15888 homo sapien	Q15893 homo sapien	ogatf70 homo sapi⇔n
SUMMARIES	ID	CONEMS	093481	C9NP07	೨೦೭೫೬೦	P72091	098605	2987T2	Garane.	055759	P77556	QBRJIO	00000000000000000000000000000000000000	Q9H4D3	Q15888	Q15893	្នុំម្នាក់ក្
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	Score		m	m	m	C4	CA	C1	C	CI	CI	C4	114	2	2	2	61
	Result No.		7	٩	4	i.c.	9	7	œ	6	10	11	12	13	14	15	16

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RESULT :

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ALIGNMENTS

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Toxoplasma gondii.
Eukaryota: Alveolata: Apicomplexa: Coccidia: Eimeriida: Sarcocystidae:
                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN-EH, CONGRAF TOTSTAIL, REA OTTER TORSRGI, AND BEVERLEY:
Lehmann T. Riackeron C P . Parmiey S.E., Kemington J.S., Dukey J.P.:
"Strain Typing of Toxoplasma gondli: Comparison of Antiqen-Coding and
Housekeeping Genes ":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         v; Indels
                                                    01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
Dihydrofolate reductase thymidylate synthase (Fragment).
Foli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     42.9%; Score 3; DB 5; Length 8;
larity 100.0%; Pred No. 6.70+05;
conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                           8 8A. 1735 8W, 33CAAAACEB133044 CRC64;
                                  A AA
                                PFI
                                                                                                                                                                                                                                                                                                                           J. Parasitol. 0:0-0(2000).
EMBL: AF249695: AAF79153.1: -.
EMBL: AF249693: AAF79150.1: -.
EMBL: AF249693: AAF79151.1: -.
EMBL: AF219694: AAF79151.1: -.
                                  PRELIMINARY;
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Pest Local Similarity
Matches o conserv
                                                                                                                                                                                                         NCBI_TaxID=5811;
                                                                                                                                                                                          Toxoplasma.
                                             09N6M5:
                              Q9N6M5
RESULT 1
                   09N6M5
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EMBL: S£4£40, AMB102120.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                    "A cytochrome of enoughd within the respiratory nitrate reductase general of Thermas Dermondalius HBB is implicated in exygen sensing under microaerophilic conditions."; Submitted (MAR-2001) to the EMBLYGENERALIGIES (ALAbases)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota, Matazoa, Chordata, Craniara, Verreksata, Puteloestomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-0cT-2000 (TrEMBLE1 15, Created)
01-0cT-2000 (TrEMBLE1 15, Last sequence update)
01-0cT-2000 (TrEMBLE1 15, Last annotation update)
Cystic fibrosis transmembrane conductance regulator (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schwarz M., Summers C., Heptinstall L., Newton C., Markham A.,
                                                                                                                                                                                                                                                                                                                                                                                Zafra O., Moreno R., Castan P., Valles C., Gava F., Caro E., Ramirez S., Berenguer J.)
                                                                                                                                                                                   Bacteria, Thermas/Celnococcus group, Celnoco i. Thermales.
Thermacee: Thermas.
NCBL_TaxID=274:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 42.9%; Score 3: DR 2: Length 10; Best Local Similarity 106.0\%; Pred, No. 3:6e+++; Matches 3; Conservative 0; Mismatches 4; Indels Matches
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                                                 01-DEC-2001 (TrEMBLrei. 19, Created)
01-DEC-2001 (TrEMBLrei. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Respiratory nitrate reductase, alpha subunit (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AJ409166; CAC69866.1; -.
NOW TER
SEQUENCE 10 AA; 1319 MW; 978E3BCB1330437A GRC64;
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FRELIMINARY.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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SEQUENCE
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10 AE

PRELIMINARY

09NP06;

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Liras P.; "Characterization of the cmcH genes of Nocardia lactandurans and Streptomyves clavuligatus encoding a functional 3'-hydroxymethylocephem dene lacinitansferase for cephanycin blosynthesis."; Gene 162:127(1995).
                                                                                                                                                                                                                                                                                                                                                  "A deletion mutation of the cystic fibrosis transmembrane conductance regulator (CFTR) locus: Delta 1507.";
Adv Exp Med Biol 290-393-398(1991)
EMRI: 964643, AAB20219.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bacteria: Firmicutes: Actinobacteria: Actinobacteridae;
Actinomycetales: Pseudonocardineae: Pseudonocardiaceae; Amycolatopsis:
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                                                                                                                            Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi:
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
51.03T.2000 (TIEMBLED. 15, Created)
61.00T.2000 (TIEMBLED 15, Last sequence update)
61.00T.2000 (TIEMBLED. 15, Last sentotation update)
63.00T.2000 (TIEMBLED. 15, Last annotation update)
63.50T.2000 (TIEMBLED. 15)
                                                                                                                                                                                                                                                                                 MEDLINE-92058253; Pubmed-1719770;
Schwarz M., Summers C., Heptinstall L., Newton C., Markham A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28.5%; Score 2; DB 2; Length 7; 100 0%; Pred. No 6 7e+05; tibe 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42.9%; Score 3; DB 4; Length 10; 100 0%; Pred. No. 3.6e+03; time 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NON_TER 10 10 SECRECT9C0544B13 CRC64; SEQUENCE 10 AA; 1107 MM; 58CFCC79C0544B13 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-FFH-1997 (ITEMHLEP). 02, Created)
01-FEB-1997 (ITEMBLED. 02, Last sequence update)
01-DEC-2001 (ITEMBLED. 19, Last annotation update)
3'-methylcophem hydroxylase (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7 AA; 746 MW; 75A1B2CDD1E681C0 CRC64;
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01-MAY-2000 (frembirel, 13, Last sequence update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=94009872; PubMed=7557411;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                     NCBI_TaxID-9506;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Transmembrane.
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2986D5;
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P72081
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"Conserved organization in the cps gene clusters for expression of
"Conserved organization in the cps gene clusters for expression of
biosynthesis acid group and the cps genes from Kitchslella pneumoniae.":
J. Bacteriol. 181:3307-2313(1999).
BMBL AFIB251: AAD300811;
BMBL AFIB251: 
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Oshima T., Alba H., Haha T., Fujita K., Hayashi K., Honjo A.,
Ikemoto K., Inada T., Itoh T., Kajihara M., Katai K., Kashimoto K.
Kimura S., Kitagawa M., Makino K., Masuda S., Miki T., Mizobuchi K.,
Mori H., Motomura K., Nakamura Y., Nashimoto H., Nishio Y., Saito N.,
Sampel G., Saki Y., Tagami H., Takemoto K., Wala C., Yamamoto Y.,
"A 718-kb DNA Sequence of Escherichia coli K-il. Genome Corresponding
to the 12-7-28-0 min Region on the Linkage Maj.",
ENAR. D90705; RAA35310.1;
Hyjothetical Frotein,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        O: Gaps
                                             Escherichia coli.
Bacteria: Proteobacteria: gamma subdivision: Enterobacteriaceae:
Escherichia.
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Escherichia,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28.6%; Score 2; DB 2; Length 8;
larity 100.0%; Pred. No. 6.76+5;
Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28.6%; Score 2; DR 2; Length 9; 100.0%; Pred. No. 6.7e+0; tive 0; Mismatches 0; Indel:
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01-MAY-2000 (TrEMPLrel 13, Last annotation update)
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                             Putative 1830 transposase (Fragment).
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Matches 2, Conservative
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Matches 2: Conservat
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                                                                                                                                                          NCBI_TaxID=562,
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FT 6
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Q9R7I2
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   SPERKRARA COOOSE SECONDE
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STRAIN=G310, AND CV. MS0; MEDLINE=G5120, Janssen D.B.; MEDLINE=G573113: Pubmed=786610; Van der Ploey J. Willemsen M. Van Hall G., Janssen D.B.; Van der Ploey J. Willemsen M. Van Hall G., Janssen D.B.; John Tanderston M. Van Hall G., Janssen D.B.; John Tanderston M. Van Hall Gariou of the halogenase gene ty insertion element IS1247 "; J. Bacteriol. 177:1348-1356(1995).

J. Bacteriol. 177:1348-1356(1995).

MON_TER 8
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Racterla, Protechacteria, alpha subdivision, Phizobiaceae group:
Hyphomicrobium group: Xanthobacter.
                                                                                                                                                                                                                                                                                                                                                 MEDINE 400 1529 PubMed*1391053; Heariques A O de Lebrastre H., Piggot P.J.; Heariques A O de Lebrastre H., Piggot P.J.; A Bacillus subtills morphogene cluster that includes spovE is browning to the Med Lepico of Escherichia coli.": EMBL: X64258; CAA45556.1: ...

BMBL: X64258; CAA45556.1: ...
                                                                                                                                                      Bacteria, Firmioutes, Barillus/Clostridium group: Bacillales:
Bacillaceae; Bacillus.
NCFL TaxID=1423;
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Best Local Similarity 100.78, Pred. No. 6.78+05;
Matches 2: Consermative 0: Mismatches 0: Inde)
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28.4%: Score 2; UB 2; Length P:
Best Local Similarity 100.0%; Pred. No. 6.7e+05;
Matches 2; Conservative 0; Mismatches 0; Indel:
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Last annotation update)
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01-NoV-1996 (TrEMBLrel. 01, Last Sequence update)
01-NoV-1996 (TrEMBLrel. 19, Last annotation update)
Haloacid dehalogenase (Fragment).
01-MAY-2000 (TrEMBLrel 13. Last sequence update)
01-MAY-2000 (TrEMBLrel. 13. Last annotation update)
Stage V spotulation protein E (Fragment).
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                                                                                                                                 Bacillus subtilis.
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                                                                                                                                                                                                                                                                                                                             STRAIN=168;
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보드보
다 - 8
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Q56759;
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Boyd E.F., Hill C.W., Rich S.M., Hartl D.L.:
"Mosaic structure of plasmids from matural populations of Escherichia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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MA W., Mac X., Jie L., Qin Z.J., Jiang W.H., Takuo R.S., Zhao G.P.;

"Clonding, sequence analysis and function analysis of the replication
order of Streptoverticilian cuestitisms infolly 12.",
Submitted (AFP-2002) to The EMR, GenBank, Tith Tarabases.

EMBL: AJ458440; CAD30324.1; -.
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Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces
NCBL_TaxID=53502;
                                                    Bacteria, Protechacteria, gamma subdimision, Enterchacteriaceae;
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Larity 100.0%; Pred. No. 6.7e+05;
Conservative 0; Mismatches 0; Indeli
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01-JUN-2002 (TrEMBLrel. 21, Greated)
01-JUN-2002 (TrEMBLrel. 21, Last sequence :pdite)
01-JUN-2002 (TrEMBLrel. 21, Last annotation upiate)
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                                                                                                                                                                                                                                                                                                                                                 Jenetics 143 1041-1100(1996).
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                  Plasmid IncFI1 E1
                                                                                                              NCBI_TaxID=562;
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Brueckmann T. Winterpacht A., Hankeln T., Schmidt E.W. Takel Brit.
Brueckmann T. Winterpacht A., Hankeln T., Schmidt E.W. Takel Brit.
Brueckmann T. Minterpacht A. 1988 13 Sequenced in DH3F project
"Human PAY" Tone RPCIPFO4K0378 1 Sequenced in Chromosome Tipls") and
Mouse (Chromosome 7) ", and "HW Pepton in Man (Chromosome III)") and
Mouse (Chromosome 7) ", and "HW Tengank" DBC databases.

British (AFF-2079) 1 The PMR, Tengank" DBC databases.

British B
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Mammalia, Eutheria: Primates: Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                        Eukaryota, Fungi, Ascomyoota, Saccharomyootina, Saccharomyeetes.
Saccharomyeetales, Saccharomyeetaceae, Saccharomyees.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. Wedler E., Scharfe M., Wambutt R.; Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (MAY-1994) to the EMBL/GenHank/DDH databases EMBL; 271169; CAA97518 2; to Sport Sono(3988; GIN1).
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U-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2001 (TrEMBLrel. 16, Last annotation update)
LIM Jonain only 1 protein (Fragment).
P87225 PRELIMINARY: PRT; 8 AA. 987225 PRELIMINARY: 047259 (TEMBLE) 12 Last sequence applate) 61 JGN-2002 (TEMBLE) 12, Last sequence applate)
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01-NOV-1996 (TrEMBLrel 01 Last sequence update)
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                                                                                                                                                                                                                                                                                                                                      Saccharomyces cerevisiae (Baker's yeast).
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GIN11 OR YLL065W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=4932;
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Lee C.-C., Yazdani A., Wehnert M., Bailey J., Guch L., Xiong M., Coolbaugh M.I., Chinault C.A., Baldini A., Lindsay E.A., Zhao Z.-Y., Caskey C.T.H.,
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                                                                                                                                                 Lee C.-C., Yazdani A., Wehnert M., Railey J., Youch L., Xiong M., Coolbaugh M.I., Chinault C.A., Raldini A., Limisay E.A., Zhao Z.-Y.,
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
(clone XPISH8A) (Fragment).
(homo sapiens (Human).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertekinta, Buteleostomi,
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                             Caskey C.T.H.;
"Isolation of chromosome specific genes by reciprocal probing of
arrayed CDNAs and cosmid libraries.";
Hum. Mol. Genet. 0.9-0(1995).
EMBL: L32069; AAA73878.1; -..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Isolation of chromosome-specific genes by reciprocal probing of arrayed cDNAs and cosmid libraries.";
Hum. Mol. Genet. 0:0-0(1995).
EMBL: L32073; AAA73883.1; -.
NON_TER 8 8
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                                                                                                                                                                                                                                                                                                                                   Query Match 28.6%; Score 2; DB 4; length 8; Best Local Similarity 100.0%; Pred. No. 6.7e++5** Matches 2; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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Joh time : 29,4167 secs
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                                                                                                                   SEQUENCE FROM N.A.
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                                                                                                                                     TISSUE-PLACENTA
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GenCore version 5.1.4\mu/.4578 Copyright (c) 1993 - 2003 Compagra Ltd.
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OM protein - protein search, using sw model

April 29, 2003, 08:30:44; Search time 16.3333 Seconds (without alignments) 17.775 Million cell updates/sec Run on:

US-09-647-749A-2 Title: Perfect score:

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Gapop 60.0 , Gapext 60.0 01.160 Scoring table:

112892 segs, 41476328 residues Searched:

0 Word size : Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

SwissProt_40:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	beschiption	P82655 lactobacill	C1	F42024			achatina	P18691 thunnus alb	P25418 libellula a	P14595 tabanus atr	P04549 periplaneta		Plinte leucophaea			980977 thunoss its					P40929 home sapies		P80677 chelyosoma	H				Adme.		∩h]am	Füli57 homo sapien	590	9918 pseud	731
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	ENTS	6 AA.	ipdate) nipdate) bbacillus s s. Lactobac	RY SEQUENCE. RC STRAIN-CB1. RX MEDITALE-1312712, FREMEd-11429463, RE The acid-stress response in Lartchacillus santranciscen RL Microbiology 147-1863-187 (2001). CC 1- INDUCTION. (VEREXPRISSED IN ANID ENVIRONMENTS. FT NON_TER 6 SEQUENCE 6 AA, 778 MR. GAA45B5B152ARGOO FROO4).	DB 1, 1		K K K	apdate) update) onent liga	Trisien v	Meighen I
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Litoria rubella (Desert tree froq).

Eukaryota: Metazoa: Chordata: Craniata: Vertebrata: Euteleusto::i.
Amphibla: Batrachia. Anura: Neobatrachia: Bufonoidea: Hylidae:
Pelodryadinae: Litoria.
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                                                                                                                                                                                              rubella.";
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                                                                                                            SEQUENCE
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            This SWISS-PROT entry is copyright. It is preserved through a collaboration between the Swiss Institute of Bloinformatics and the EMBL outstation the European Bloinformatics. There is no restrictions on its use by non-profit institutions as long as its content is in no way mobilied and this statement is not removed. Rouge by and for commercial waitings in purposed and this statement is not removed. Rouge by and for commercial waitings in a number of send an email to inconsedable shock).
              -!- CATALYTIC ACTIVITY: ATP + an acid + protecus = AMP + diphosphate +
                 - CATADITIC ACCOUNT TAILORS HER.

an acyl-protein thioleseer.

PATHWAY: Bioluminescent fatty acid reduct... System; second step.

- PATHWAY: Bioluminescent fatty acid reduct... System; second step.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebinta, Euteleostomi,
Amphibia: Batrachia, Anura, Neobatrachia, Bufonnidea, Hylidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE-Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowne J.H., Gao C.,
Tyler M.J., Wallace J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -i-TISSUE SPECIFICITY: SECKELED BY THE SKIN LIBRAL GLANDS.-i-MASS SPECIFOMETRY: MW=655; METHOD*FAB.
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1.1e+05
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S AA: 656 MW: 71AYCYCB10300000 CR: 44;
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                                                                                                                                                                                                                                                                                              Ouery Match
28.6%, Score 2, DB 1, (w)
Best Local Similarity 100.0%, Pred. No. 1.1e+05;
Matches 2, Conservative 0; Mismatches
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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RE31_LITRU STANDARD; PRT; 5 AA. PR2072; 15-508-200-2002 (Rel. 41, Created) 15-5008-2002 (Rel. 41, Last sequence update) 15-5008-2002 (Rel. 41, Last sequence update) Rubellidin 3.1.
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Luminescence; Ligase.
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MoD RES 5 5
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NCBI_TaxID+104895;
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ses 2, Conserv
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P82073;
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ID RE32_LI
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MEDLINE=$5666991 PubMed-7970891, Maule A.G., Shaw C., Bowman D.W.; Maule A.G., Shaw C., Bowman D.W.; Maule A.G., Shaw C., Bowman D.W.; The EMRERAIDS INCREDUCE insurance in the fire living increduce. Faragrafulus redivitus (Nematoda, Pharbitida).") Parasitology 109:381-356(1994).

FIRETOR HAE PERECTS ON MISCLE TENSION.

FIRETOR FREDIVITY. FOUND IN THE NEWER CORDS AND A VARIETY OF GANGLIA FARTICULARLY IN THE ANTEROR PROJECTS.
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MEDLINE=93324431; PubMed~8332542;
Cowdon C., Stretton A.O.W.;
Cowdon C., Stretton A.O.W.;
TaP2, an Ascaris neuropeptide: isolation, sequence, and bivactivity.
Paptides 14-423-430(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota: Metazoa: Nematoda: Chrimadorea: Ascaridida. Ascaridoidea:
TISSUP-SPID secretion:
Wakhite P.A., Bowie J.H., Tyler M.J., Wallace J.C.,
"Poylid-s from Sovie J.H., Tyler M.J., Wallace J.C.,
Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                     ust. J. Cham. 52:639-645(1999).
!- PGNCYION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBICITA
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                                                                                                                                                                                                                                        -:- TISSTE SPECIFICITY: SECRETED BY THE SKIN DORSAL GLANDS
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SEQUENCE 7 AA; 992 MW: 64946779RFRITE359 CP264;
                                                                                                                                                                                                                                                                                                 5 AA; 570 MW: 71A9090'852A0000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-701 1993 (Fel 26, Created)
01-701-1993 (Rel. 26, Last Sequence update)
1-101-1996 (Rel. 23, Last aumotation update)
FMRFamide-like neuropeptide AF2.
                                                                                                                                                                                                                                                                                                                                          28.6%: Sic. 100.0%; Pred. No. ...
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Best Local Similarity 100.0%;
2: Conservative
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Matches 2; Conservative
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NCBI_TaxID=6253, 6233;
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01-NOV-1990 (Rel. 16, Last sequence update)
01-NOV-1990 (Rel. 16, Last sequence update)
01-NOV-1990 (Rel. 16, Last annotation update)
01-NOV-1990 (Rel. 16, Last annotation update)
Anglotensin-converting enzyme inhibitor.
Thunnus albacares (Vellowfin tuna) (Neothunus macropterus).
Thunnus albacares (Vellowfin tuna) (Neother tar Euteleostei)
Acanthomorpha, Neotherygil, Teleostei, Euteleostei, Neoteleostei,
Acanthomorpha, Aranhopherygil, Percomorpha, Percoformas, Scombridae; Thunnus.
                                                                                                                                                                                                                                                                  Eukaryota, Metazoa, Mollusca, Gastropoda, Fulumunata, Stylommatophota,
Achatinacea, Achatinidae, Achatina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kohama Y | Marsumoro S | Oka H | Teramoto T | Fashe M | Mimura T | Toolation of angiotensin-converting enzyme inhibitor from tuna
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SEQUENCE 9 AA; 953 MW. 6AA843734051F1R7 PPO44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7 AMIDATION.
965 MW, PREZISERFORTSZPIS ZN 54,
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100 0%: Pred No 1 1++09
ive 0; Mismatches 0.
                                                                                                                 01-JUN-1994 (Rel. 29, Greated)
01-JUN-1994 (Rel. 29, Last Sequence update)
01-OCT-1994 (Rel. 30, Last annotation update)
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Kohama Y., Mafsumoto S., Oka H., Teramoto T
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. w.:
Five 0; Mismatches
                                                                                                                                                                                                                                            Achatina fulica (Giant African snail)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-93265912: PubMed-8495720;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hers Lett 323 luq-luu,
PIEN S3244, S33244
Weit FFFFSE, Andatist,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2) Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
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                                                                  SIANDARD,
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=6530;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.
                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE.
TISSUE-Ganglion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE-Muscle;
                                                           WWA3_ACHFU
P35921;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACI_THUAL
P18691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 KE 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - X
- X
- X
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P25418;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 KE 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 IK 2
                                                                                                                                                                                                                WWamide-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ระบูกหลาก
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RESULT 6
WWA3_ACHFU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 8
AKH_LIBAU
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TISSUE=Corpora cardiaca,
MEDLINE=90046758; pubmed=2814485;
Jaffe H. Baina A.K., Piley C.K.,
Jaffe H. Paina A.K., Piley C.K.,
"Filminy structure of the rest periods burnings with adjrithetic and hyperplants structure of the rest periods the corporal cardiaca of horse files (Diptera)."
Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sars
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tabanus atratus (Horse fiy).
Fukaryota: Matazoa: Arthropoda: Mandibulata; Pancrustacea: Hexapoda:
Insectoa: Pterygota: Nechtefa: Endopterygota: Diptera: Brachycera:
Tabanomorpha: Tabanidae: Tabanus.
NGFLTaxID=7207;
                                                                                                                                           Hetapolas
01-MAY-1992 (Rel. 22. Created)
01-FBB-1994 (Rel. 28. Last sequence update)
01-FBB-1994 (Rel. 28. Last sequence update)
Adipokinetic hormone (AKH).
Adipokinetic hormone (AKH).
Edibelials auripensis (Skimmer dragonfly).
Edkaryota Metazoa Akhrogoda Mandiboliata, Fanctistarea metazoa Antagoda Insecta; Pterygota: Paladesptera: Odonata Antagota Libelia: Gard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "The putative appostral poptide of the adiportnetic/red-pidment-concentrating hormone family isolated and sequenced from a dragonfly.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JAN-1990 (Rel. 13, Created)
01-FBR-1994 (Rel. 28, Last Sequence update)
01-FBR-1994 (Fel. 28, Last sequence update)
01-FBR-1994 (Fel. 28, Last semintation update)
Adiporting inclination (AKE) (Dipperan corpora cardiaca factor I)
(DCC I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 28.68: Score 2: DB 1; Length 8; Best Local Similarity 100.0%; Pred. No. 1.18+05; Matches 2; Conservative 6; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PYRROLIDONE CARBOXYLIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8 AA: 978 MW: 8565A771A9C452D6 CRC64;
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                                                                                                                                                                                                                                                                                                                        TISSUE-Corpora cardiaca;
MEDLINE-90359055; Pubmed-2390213;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          InterFro, IPR002047; AKH.
PROSITE, PS00256: AKH: 1.
Heuropeptlide: Amidation; Flight.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PROSITE: PS00256: AKH; 1.
Neuropeptide; Amidation; Flight.
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InterPro; IPR002047; AKH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                            SECUENCE, AND SYNTHESIS.
                                                                                                                                                                                                                                   NCBI_TaxID=6966;
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P14595;
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SEQUENCE
                                                                                                                                                                                                        Libellula.
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      WERRAND COCOCORRAND WAS COLORED TO THE PROPERTY OF COCOCOCORRAND WAS COLORED TO THE PROPERTY OF COLORE
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1. FUNCTION: HYPERTREHALOSED THE THE WHYDIXMPH (PEHALOSE IS THE ELEVATE THE LEUR IN THE HEMOLYMPH (PEHALOSE IS THE MAJOR CARBOHYDARE IN THE HEMOLYMPH (PEHALOSE).
                                                                                                                                                                                                                                                                                                    13-403-1987 (Rel. 05, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Hypertichalosaemic factor if Neuropeptide Nift (Feriplanetin CC 2)
Feriplaneta americana (American cockroach)
Periplaneta americana (American cockroach)
Leptinotarisa decembiracia (Periplaneti Periplaneti II)
Entra orientalisa (Oriental cockroach)
Enkaryota: Metazoa: Arthropoda: Mandibulata: inncrustacea: Hexapoda: Insecta: Periplaneti Cockroach)
Entroidea: biaticidae: Periplaneta: Orthopteroidea: Sictyoptera: Blattaria:
WCBL_TaxID=E978, 7539, 6976;
                                                                                                             Saps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SPECIES-P.americana:
MEDLINE-84238179; PubMed-6591205;
Scarborough R.M., Jamieson G.C., Kalish F., **amer S.I., McEnroe G.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The metabolic neuropeptides of the corpus cardiacum from the potato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Richart K.L. Jr.:
"Structures of two cockroach neuropeptides acciqued by fast atom
bombardment mass spectrometry.";
Bicchem. Ricphys Res. Commun 124-250-359(1044);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-8594653); PsteMod-6548629;
Witten J.L., Schaffer M.H., O'Shea M., GOSK ''., Hemling M.E.,
Riochart K.L. Jr.;
                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Isolation and primary structure of two peptides with cardioacculatory and hyperalycemic activity, from the corpora cardiaca of Periphaneta americana", proc. Natl. Acad. Sci. U.S.A. 81:5575-5579(1994).
   PYRROLIDONE CAPHOXYLIC ACID.
                                                                                                             "; lndels
                                                                            Length 8;
1 1 FIRECLE...
8 8 AMIDATION.
8 AA; 949 MW: 80786771A9D1A736 CRO54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             beetle and the American cockroach are idential.":
Peptides 10:1287-1289(1989).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPECIES-L.decemlineata: TISSUE-Corpora card.ora;
MEDLINE-90160053; Pubmed~2576128;
                                                                          28.6%, Score 2, DB 1 in Liarity 100.0%; Pred, No. 1.1e+4. Conservative 0; Mismatches "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SPECIES-B.orientalis; TISSUE-Corpora cardiada:
MEDLINE-90253659; PubMed-2340112;
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                                                        wery Match
Best Local Similarity
Tobes 2: Conserva
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PIP: S08995; S08996.
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   MOD_RES
MOD_RES
SEQUENCE
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01-TAN-1990 (Rel. 13, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annitation update)
Leukopyrokinin (LPK) (LEM-PK).
Leucophaea maderae (Madeira cockroach).
Leucophaea maderae (Madeira cockroach).
Leucophaea maderae, Hexapoda;
Insecta, Pierygota; Neoptera, Orthopteroidea, Dictyoptera: Blattaria:
Riaberoidea, Blaberidae, Leucophaea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saps
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01-AUG-1991 (Rel. 19, Last sequence applate)
101-AUG-1991 (Rel. 19, Last annotation update)
101-AUG-1991 (Rel. 19, Last annotation update)
101-AUG-1991 (Rel. 19, Last annotation update)
101-AUG-1991 (Rel. 1991 (Augusta) Augusta, Augu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schoofs I. Holman G.M. Hayes T.K. Nachman R.J. de Loot A.:
"Isolation, identification and synthesis of locustamyotropin II. an additional neuropephide of Locusta migratoria. Member of the repealomyotropic peptide family.":
Insect Riochem. 20.4794-484(1940).
-i. FUNCTION: MEDIATES VISCERAL MUSCLE CONTRACTILE ACTIVITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
                                                                                                                                                                        PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                  28.6%: Score 2: DB 1: Length 8: 100.0%; Pred: No. 1.1e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 28.6%; Score 2; DB 1; Length 8; Best Local Similarity 100.0%; Pred. No. 1.1e-05;
                                                                                                                                                                                                                                                 86745771A9D1A736 CRC64:
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                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. ....
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                                                                                                                                                                                                              AMIDATION.
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                                                                                                                                                                            MOD_RES 1 1 1
MOD_RES 8 8 8
SEQUENCE 8 AA; 1005 MW;
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                                                                                                                                                                                                                                                                                                                                                                                               2; Conservative
                                                             INTELPRO: 1PR002047; AKH. PROSITE; PS00256; AKH. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE=Corpora cardiaca:
Schoofs L., Holman G M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STANDARD
                                                                                                                                         Neuropeptide: Amidation.
PIR; B44960; B44960.
FIR; B49823; B49823.
                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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P22396;
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P13049;
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4 FT 5
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LPK_LEUMA
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MEDINE-2041310; PubMed-10952880; Dircksen H., Burdzik S., Sauter A., Keller P.; The correction to the highly "Two creatines and the movel ortapeptide order or any fish brokened ilmosus ilmosus ilmosus of the crayfish brokenedtes ilmosus ilmosus of the terminal neuropeptides originating together in neuropes of the terminal
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**MISCELLANBOUS: AN ANALOG WITHOUT THE N-TERMINAL FOA RESIDUE WAS SYMPHESIZED AND FOUND TO EXHIBIT SPEATER ACTIVITY (144%) THAN THE PARRIMY BURROPEDTIDE. THE PORTION OF THE SEQUENCE OF LEK MOST CRITICAL FOR THE MYOTROPIC PROPERTIES IS LIMITED TO THE PROPERTIES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Metazoa, Ařthřopoda, Mandibulata, Pacifustacea, Crustacea,
Malacosfraca Eumala-Nestraca, Eucafida, Delapoda Plebogemata,
Astacidea, Astaccidea, Cambaridae, Orconocios, Nebogemata,
NCBI_TaxID+28479;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Subs
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FUNCTION: MYOTROPIC PEPTIDE, ENHANCES BOTH THE PREQUENCY AND
AMPLITUDE OF SPONTANEOUS HINDGUT CONTRACTIONS. IT IS SYNTHESIZED
                                                                                                                                                                                                                                                                                                                       MEDINE-87052651; FubMed-2877794;
Holman G.M., Cook B.J., Nachman R.J.;
"Primary structure and synthesis of a blocked ejettopic
neuropeptide, solated from the bookroach, Leonophaea maderael",
Comp. Blockem, Physici, 85c.219-224(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PYRROLIDONE CARROXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      e Indels
Nachman R.J., Holman G.M., Gock B.J.; "Active fragments and analogs of the insect serropeptide leucopyroklinis structure-function studies.": Biochem. Biophys. Res. Commun. 137:936-942(199.).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1, Longth 8, b. 1.18+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 8 AMIDATION.
8 AA; 965 MW: 87086181A90DDDAA9 CR044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8 AA; 949 MW; 92341771A9D5A1R6 CRC+4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- SIMILARITY: BELONGS TO THE PYROKININ FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-0CI-2001 (км) 40, стеятед)
16-0CT-2001 (Rel. 40, Last sequence update)
16-0CT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BY ABDOMINAL GANGLIONIC NEURONS.
1- MASS SPECTROMETRY: MW-904.8; METHOD-FAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. No. 1.1
ive C, Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Orcomyotropin (OMT).
Orconectes limosus (Spinycheek crayfish).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28.6%; Score 2;
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PROSITE: PS00539: PYROKININ: 1.
Neuropeptide: Amidation: Pyrokinin.
MOD_RES 8 AMIDATION: Pyrokinin.
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                                                                                                                                                                                                                                        SEQUENCE, AND SYNTHESIS.
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Best Local Similarity
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P82455;
                                                                                                                                                                                                                                                                                          TISSUE-Head
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 TS 3
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ORMY_ORCL1
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01-071-1994 (Fel. %) Last Sequence update)
15-080-1998 (Fel. %) Last annotation update)
Cardioactre peptide (CAP).
Mandata maenas (Common Manche Crab),
Mandata sexta (Tokacco hawkment) (Tokacco horinworm),
Penchric molitor (Yollow mealworm), and
Speciation and Southern armyorm).
Eukaryota, Melazoa, Arthropoda; Mandibulata; Panorustacoa; Cissiacea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Stangier J. Hilbich C. Beyreuther K. Keller R.; "Unsual cardioactive peptide (CCAP) from pericardial organs of the shore crab Carcinus meanes.": Proc. Natl. Acad. Sci. U.S.A. 84:575-579(1987).
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Cheung C.C., Loi P.K., Sylwester A.W., Lee T.D., Tublitz N.J.:
Primary Structure of a cardioactive neuropeptide from the tobacco
hawkmorth, Mandura sexta ".
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01-NOV-1997 (Rel. 35, Last sequence update)
15-Jun-1994 (Rel. 38, Last annotation update)
15-Jun-1994 (Rel. 39, Last annotation update)
17-Jun-1994 (Rel. 39, Lest annotation update)
Thunnus obesus (Rigeye tunns)
Eukäryota: Metazoa: Chordata: Craniata: Vertebrata, Estereostomi:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Malacostraca, Eumalacostraca: Encarida, Perapoda, Piecovemata;
Brachyura; Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBL_TaxID=6755, 7130, 7067, 37547;
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1 10+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             M. Smatches
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Lattiy 300 2%; Prad No
Cusservativo 6; Mismatch
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                                                                                          PRT;
                                                                                                                                                                               01-OCT-1994 (Rel. 30, Created)
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FIR, 627233, 627233.
Neuropeptide, Amidation.
                                                                                          STANDARD:
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Watches 2, Cosserv
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PR0975;
                                                                                          CCAP_CARMA
P38556;
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SEQUENCE
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RESULT 14
CCAP_CARMA
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-1- FUNCTION: THIS PROTEIN IS ONE OF THE NUCLEAR-CODED POLYPEPTIDE CHAINS OF CYTOCHROME C OXIDASE, THE TERMINA. OXIDASE IN MITOCHORNAL ELECTRON IRANSPORT.

-1- CATALYTIC ACTIVITY: 4 FERIOSPORT.

-1- CATALYTIC ACTIVITY: 4 FERIOSPORT.

-1- SIBILARITY: ALCOHORNAL LOCATION: Mitochondrial inner numbrane.

-1- SIBILARITY: HELDINGS TO THE CYTOCHROME C (XIDASE VIA FAMILY. InterPro: IPE00149: COX6A. PRRTIAL.

PROSITE: PS01329: COX6A: PARTIAL.

PROSITE: PS01329: COX6A: PARTIAL.

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NON_TER 5
Actinopteryqil; Neopterygii; Teleostel; Euteir stel; Neoteleostel; Acanthomorpha; Acanthopterygii; Percomorpha; Perciformes; Scombroidel; Scombridae; Thunnus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                     "The subunit structure of cytochrome-c oxidase from tuna heart and
                                                                                                                                  Arnold S., Lee I., Kim M., Song E., Linder D., Lottspeich F.,
Kadenbach B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          o Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 28.6%; Score 2; DB 1; length 9; Best Local Similarity 100.0%; Pred. No. 1.1e+05; Matches 2; Conservative 0; Mismatches 0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          9 AA: 1136 MW; 62E07249480776DE (04064)
                                                                                                  MEDLINE-97454291 PubMed-9310366; MEDLINE-97454291 FubMed-931036
                                                    NCB1_TaxID=8241;
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NON_TER
SEQUENCE
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Search completed: April 29, 2003, 08:52:29 Job time: 17.3333 secs

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GenCore version 5.1.4 _{\rm L} f \le 45.78 Copyright (c) 1993 - 2002 Compute Little.
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OM protein - protein search, using sw model

Apili 29, 1803, 08:45.15 , Search the 29.1007 Seconds (Without Alignments) 23.672 Million cell updates/sec Run on:

US-09 647-749A-2 1 IKEYFTS 7 Title. Perfect score: Sequence: OLIGO Gapop 60.0 , Gapext 60.0 scoring table.

Word size :

287224 seqs, 96134422 residues

Searched.

Total number of hits satisfying chosen parameters.

1100

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

Database :

ptR_73.* 1. ptrl:* 2. pir2:* 3. pir4:*

Pred. No. is the number of results predicted by chance to have a source greater than or equal to the score of the result being printed, and is derived by analysis of the total so or distribution

SUMMARIES

		*			SUMMARIES	
Result.	Score	Cuery Match	Length	DB	. di	Description
-	~	42.4	6	: ~	A24244	adipokinetic hormo
- 04	· m	42.3	10	7	6144	polygalacturonase
~	71	∵. 83	S	\sim 1	B37325	pap fimbrial requi
-7	2		5	~	T027	neavy chai
en	7		ur.	a	064	-cell receptor
æ	C4	30.0	5	C4	(1)	. S. S.
r	2		ic.	~1	-	4.5K structural
æ	CI		u)	C	448	M 10
6	CI		5	C.	-1	5K structural pr
10	CI		ď.	C4	111	Ţ.,
11	CI	38.6	9	C1	A31263	dihydrofolate redu
31	1.4		r -	, ,	315118	alucese isomerase
13	CI		r.	C 4	F30254	
14	C1	9. 82	L.	C4	533244	neuromodulatory pe
15	CI		r·	C4	PC1316	large granule L2 c
16	C1		L.	C4	A28671	peptidylglycine mo
17	7	28.6	7	~	A38081	amine oxidase (cop
18	CI	3.9.4	r	7.5	A59725	wirotoxir - dostro
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21	7	58.6	α	CI	\$10596	adipokinetic hormo
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23	C)		x	-1	7	
24	Cŧ	28.6	œ	C#	566	adipokinetic hormo
25	5	28.6	80	CI	5531	
5.6	CI		œ	(1	5862	adipokinetic hormo
27	C1		ထ	C4	~4	leghemoglobin III
28	64		œ	C4	3157	Ö
67	CI	28.6	သ	Cŧ	32	hypothetical prote

5 Q.

Tariant surface gi leucopyrokinin - M homeoric profein u alcohol debydrogen serum albumin - do fibroblast growth cytochrome F45c AL MATTRASSPORTING A MAC class I bistoc phyllocerulean translation elocga leader peptide - P bradykinin-like pe Threb bradykinin- kradykinin-like pe bradykinin-like pe	RESULT 1 A4424 adjournette hormone - bollworm Nalternate names: Hez-AKH adjournette names: Hez-AKH adjournette names: Hez-AKH c.Species. Hellothis zea (bollworm, corn ediworm, tomato fruitworm) C.Date: 31-Mar-1988 #sequence_revision 23-Mar-1995 #text_change Nict-1997 B.Caffee H. Raina A.K.; Piley, C.T.; Fraser, P.A.; Holman G.M.; Wagher, P.M.; Piley B.Caffee H. Raina A.K.; Piley Niction 185, 622 628, 1986 #text_change Niction Nict-1997 A.Title: Isolation and primary structure of a peptide from the corpora condince of He A.Recession: A24244; MIID:86186794; PMID:3964263 A.Recession: A24244; MIID:86186794; PMID:3964263 A.Recession: A24244; MIID:86186794; PMID:3964263 A.Recession: A24244; MIID:86186794; PMID:3964263 C.Specifanily: A342625 hottle hormone	4.2: Length 9: 2.8e+05; 3.	8 pg	b 2) Length 10; 1.1e+03; es 0; Indels 3: Gaps 0;
22 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	orm Lworm, coin earwo Lregislon 23-Mar- ey, C.T.; Praser, 115, 622 628, 1 Tructure of a MIID:86186794; PMI cormone Lend cospect car attack; end (512)	42,4%; Score %; DR 2. 100,0%; Pred, No. 2.8R 150e C. Mismatchess		42 4%; Somm %; DB 2) 100.0%; Pred. No. 1.1e- ive O: Mismatches
	hormone - bollwe names: Heal (bu) kar-1988 #sequence A244	ry Match t Local Similarity 100 ches 3: Conserration 5 FTS 7 111 4 FTS 6	coase (EC 3.2. spergillus Sp. D61440 mark.t. himse Aspergil number: A61410 belim har; refer protein 1.10 STS.	similarity 2. Oonservat
ರಟಬಹಕಾಗಿ, ಐಳಂದಾವವವವ ರಟಬಹಕಾಗಿ, ಐಳಂದಾವವವವು	RESULT 1 A24244 adjpokinetic N. Shecies. B C. Date: 31-N C. Accession: Bicchem. Bic A. Title: Isc A. Reference A. Rocession: A. Reference A. Residues: C. Experient A. Residues: B. C. Experient A. Rocession: A. Ro	Ouery Match Rest Local Matches Oy S FT Db 4 FT	PESULT 2 D6149alactur D6149alactur D78acies: R C78ae-8 n77-8 C78ae-8 n77	query Mato Best Local Matches

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34.5K structural protein - Leuconostoc oenos phage P54 (fragment)
C;Species: Leuconostoc oenos phage P54
C;Sate, 31.3ar.1933 *sequence_revision 12.May.1998 *text_nhange 21.May.1949
C;Accession: F44817
B;Arendt, E.K.: Lonvaud, A.: Hammes, W.P.
J; Gen M;Archiol. 137 2134-2130 1943
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817: Multi-q2ngGn33: PMID-1748868
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7. Stariotural protein - Leuromostor cenos phage P2t11-15 (fragment)
7. Starios Leuromostor enes phase P2t11-15
7. Starios Leuromostor enes phase P2t11-15
7. Date: 31-Mar-1993 #sequence_rewision 22-May-1998 #text_change 22-May-1998
                                                                                                                                                          34.5K structural protein - Leuronostor cenos phage P32 (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 28.6%; Score 2; DB 2; Length 5; Length 5; Length 5; Conscripting 0; Mismatches 0; Telois es 2; Conscripting 0; Mismatches 0; Telois
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A,Residues: 1-5 <ARE>
A,Note: segmence extracted from NORI backhone (NCPIP:70342)
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A;Residues: 1-5 <ARE>
A,Note: sequence extraple(from hrp! backbase (hrpph=70332)
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A,Residues: 1-5 <ARE>
A,Note: sequence extracted from NCBI backbone (NCBIP-70:35)
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C)Accessions 184487.
E)Arendt E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Mirrobiol. 137, 2135-2139, 1991
A.Title: Lysogeny in Leuconostoc Oenos.
A.Ferferbiol. Gamber. A44817, MRID 92785072, FMID 1748868
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                                                                                                                                                                R:Braaten, B.A.: Blyn, L.B.: Skinner, B.S.: Low, P.A.
J. Bacteriol. 173, 1789-1800, 1991
A;Title: Evidence for a methylation-blocking factor (mbf) locus involved in pap pilus ex
A:Reference number: A33325; MUID:91154136; PMID:1-71857
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R.Yanada M.; Wasserman, P.; Feichard, B.A., Shan, S., Caton, A.J., Elvera, G. J. Exp. Med. 173, 395-407, 1991.
A.Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A.Reference number: Pr0222; MUD:91108337; PMID:18/90102.
A.Reference number: Pr0222; MUD:91108337; PMID:18/90102.
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A;Tille: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:171.558
A;Recession: PT0504; MUID:91277601; PMID:171.558
A;Recession: FT0544
A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: LF5 cFEEA
A;Residues: LF5 cFEEA
A;Residues: LF5 cFEEA
A;Residues: T5 cFEEA
C;Reywords: T-cell receptor
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C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 *sequence_revision 30-Sep-1993 *text_change 16-Aug-1996
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C.Date: 1-ful-1992 #sequence_revision [7-7,].1992 #th.z<sup>1</sup>_channe 30 May 1993
C.Accession: PT0644
R.Feeney, A.J.
                                                                                    C:Species: Eschérichia coli
C:Date: 11-Sep-1992 #sequence_revision 11-Sep-1997 #text_change 23-Mar-1993
C:Accession: B37325
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P10644
Thoball receptor beta chain V-D-J requon (111-13) in mouse (fragment)
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pap fimbrial regulatory protein papI · Escherich... coli (fragment)
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A.Residues: 1-5 < YAMA
A.Residues: 1-5 < Ource: B lymphocyte
C.Keywords: heterotetramer; immunoylobull::
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Exp. Med, 174, 115-124, 1991
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Matches 2; Conserv
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A:Molecule type: DNA
A:Residues: 1-5 <BRA>
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R:Yanada M.; Wasserr
J. Exp. Med. 173, 399
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Sdep .a

o: Indels

y. 150

RESULT 9

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discose isomerase - Thermognaerobacterium saccharolyticum (fraument)
C.Species. Thermognaerobacterium saccharolyticum
C.Species. Thermognaerobacterium saccharolyticum
C.Species. 128-1948 #sognaro_rorizion |2-free.jqua #foxt_rhange |T.Aprilong
C.Species. 273-565-571
Blochem. J. 273, 565-571
Blochem. J. 273, 565-571
Blochem. J. 273, 565-571
A.Ancession. S17976
A.Reference number: 815119; Mg10:91144536; PMID:1996956
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E.Minskitt, E., IM-Ba, T., Manacka, Y., Febayashi, M., Memits, F.

FPBS Lett, 323, 104-108, 1993

Affilte: Whamiderl, 2 and 3; novel neuromodulatory peptides isolated from sandila of Reference number: $33244; MUD:94265912; PMID:8495720

A.Rocession: $33244; MUD:94265912; PMID:8495720
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C.Species: cryza sativa (rice)
C.Date: G3Feb-1994 #sequence_tevision 03 Feb-1994 #text_change 23-Mai-1945
C.Datession: ps0254
R.Tsugita, A.
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C.Species: Achatina fulka (glant African snail)
C.Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change_14-71.-1947
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A;Wckeule type: protein
A;Pssidues: 1-7 <150
A;Experimental source: leaf, chloroplast, strain Nihonbare
A;Note: molecular weight 18K, pl 4.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0
Matches 2; Conservative
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A:Reference number: PS0206
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A.Molecule type: protein
A.Residues: 1-7 <MIN>
1 TS 2
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SIGNAT

alcohol dehydrogenase (EC 1.1.1.1) chi - horse (fraument)

C.Spectes: Bquus caballus (domestic horse)

C.Date: 28-Feb-1990 #sequence_revision 29-Feb-1990 #text_change 31-Jun-1997

C.Accession: 802617

R.Fairwell, T. Julia, F. Kaiser, P.: Holmquist, F.: Pares, X.: Vallee, B.L.: Journvail

R.Fairwell, T. Anila, F.: Kaiser, P.: Holmquist, F.: Pares, X.: Vallee, B.L.: Journvail

R.Fairwell, T. Anila, F.: Raiser, P.: Holmquist, F.: Pares, A.: Vallee, B.L.: Journvail

A.Title: Acceptated N.terminal structures of class :II alochol dehydrogenases. Difference

A.Reference number: 802617; MuID:88005160; PMID:367405
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A31268
A) Thymidylate synthase (EC 1.1.1.45) - Flashedrum talcharum
C)Species Basmedium talcharum
C)Species Basmedium talcharum
C)Species Basmedium talcharum
C)Accession: A31263
A) Accession: A31263
A) Title: Evidence that a point mutation in dihydrum talcharum
A) Reference number: A94217; MUID: 89057886; PMID: 2044149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A)Molecule type: protein
A)Residues: 1-6 <FMI>
C:Superfamily: alcohol dehydrogenase; long-chain alvohol dehydrogenase homology
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A,Status: not compared with conceptual translation
A,Molecule type: DNA
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Conservative

Best Local Similarity Matches 2; Conserva

Query Match

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RESULT 11

1 IK 2

3 TS 4

5 a

5 IS 7

Residues: 1-6 <PET>

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Up 2 KE 3

Up 2 KE 3

BESULI 15

PC1316

PC1316

PC1316

C.Dactanule L3 chain - horseshoe crab (Tachypieus tridentatus) (fragment)

C.Species: Tachypieus tridentatus

C.Accession: PC1316

A.Title: Separation of large and small granules from horseshoe crab (Tachypieus tridentatus)

A.Peference number: PC1309; MUJD:94110249; PMID-82M:718

A.Rocession: PC1316

A.Roce
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GenCore version 5.1.4_F5_4578 Copyright (c) 1993 - 2003 Computer Ltd.
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OM protein - protein search, using sw model

April 25, 2003, 08.30:19 , Search time 58.9167 Seconds Run on.

(without alignments) 15.832 Million cell updates/sec

US-09-647-749A-2

1 IKEYFTS 7 **Title:** Ferfect score: Sequence: Scoring table:

ublGo Gapop 60.0 , Gapext 60.0

408470 seqs, lasabbeab residues

Searched:

Word size :

fotal number of hits satisfying chosen parameters:

174064

Minimum DB seq length: 0 Maximum DB seq length: 10

Post-processing: Listing first 45 summaries

Database :

A_Geneseq_101002:* A_Geneseq_101002:* SIDS_AGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score		% Ouery Match Length DB		qı	Description
: -		100.0		1 3	AAY43696	HIV-1 aplie induce
· (1	**	(-)	ſ	C1	AAB55781	PDZ motif sequence
~	47	57.1	S	C1	AAB57513	Glycophorin C C-te
7	4	57.1	S	C1	AAB58023	Glycophorin C C-te
L ⁿ	7	1.7.1	w	C 1	AA.B55782	FDi motif sequence
æ	4	57.1	٩	22	AAB57614	Glycophorin ć C-te
7	4	57.1	9	C4	AAB58024	Glycopheria C C-te
œ	4	57.1	7	-4 (4	AAB55783	FD2 motif sequence
Ţ	4	5.7	1	C1	AAB57615	Glycophorin d c-te
2		٤,7	7	2.2	AAB58025	Glycophorin C C-te

iseque	Shortin C O-t	le fradment	anic peptide	itor	acid seque	LA-A2	scadherin ce	53 DR 3a motif	SP immunogen	580107	59F1D7 related	erived imm	rotocadherin ce	umar.ised	man complement	SPICT relate	SPID related	D related	Plr related	Sarin rela	PIII related	SSPIET relate	47 class II p	Copeptide iseful :	1. 1. 1. 1. 1.	motif seque	ermina	ophorin C	1	shorit. 🤔		death prote	D element in estro
B5578	AABS	AAR5922	AAR9754	AAT5336	AAY4018	AAY2670	AAY63	AAG89	AAB76	ABJ01	ABJ01	AAR	AAY63	AME12	AAG45	ABJ01	ARTO!	ABJ01	ABJ01	AHJ@1	ABJ01	ABJUL	AAY 82	AAYSB	AABSS	AAB5578	AAB5748	AAB5761	AAB5783	AABTEC	£AU7891	44069	AAR0737
4 57.1 8	7.	1	7.1	7.1	7.1	e1 F:	7.1	7.1	7.1	7.1	٦.		7.1 1		7.1 1	7.1 1	٠. ١	7.1 1	7.1 1	7.1	7.1 1	7.1	5.5	5.9	42.4	42.9	42.	42	7	: 4 - 7	-,-	42.3	
п. П.	4 F	1.4	15	16	17	0) e4	19	2.0	21	22	23	24	25	2.6	27	7.8	50	3.0	3.1	3.2	3.3	3.4	3.5	3.6	3.7	χυ ×	3.6	40	-,	() -#	~		u: T

SIMBMEDIE

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Mentinginal to protective discuss, HIV-1, ecretope protein; uplicitively, near a ressent dial ocl. chemokine, that interferent neutronal rail leath, every final ocl. chemokine, that interfere demential deptession, psychologis, opportunistic infection: morrotexicity; infantary percelegical disease, multiple scherosis, tropical spastic paraparegis; Alzheimer's disease, multiple scherosis;
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                                                                                                       AAY43636 standard, Pertide, 7 AA.
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FESULT 1
                                                             AAY49696
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Synthetic.

W09951254-A1.

Pr-App-1999; 14-0CT-1999.

GRITS-DARGRISE. 06-APP-1998:

(ADIM-) ADVANCED INMUNIT INC. Pert C, Ruff M;

WPI; 1944-633645/54.

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responses, e.g. inflammation, cancer
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000008-0196460.
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Best Local Similarity
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                                                                     AAY49695 and AAY49866 represent pertites which inhibit HIV 1 Jpl20 induced neuronal cell death. Pharmaceutical compositions containing the peptides are useful for treating symptoms caused by meatonal cell less such conditions especially associated with HIV infection include encephalopathies, neuropathies, memory loss, dementa, depression psychosis and opportunistic infections. The populage act as antagonists of apt20-mediated neurotoxicity and subsequent neuronal degeneration. This enables therapeutic treatment of HIV intestion and other inflammatory neurological diseases, including multiple sclerosis, tropical spastic paraparesis and Alzhelmer's itsease.
New peptides useful for inhibiting human [mman-deficiency virus type 1 (HIV-1) gp120 induced neuronal cell death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modulating a biological function of an audothelial cell or
hematopoletic cell, useful for treating autoimmune diseases and
infectious diseases, by administering an autogonist that inhibits
binding between a PDZ protein and a PL protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a new meth d for modulating a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hematogeletic ell, FDE, FL, autoimmune liseuse, inflammation, allergy, asthma; multiple sclerosis; cancer, infection.
                                                                                                                                                                                                                                                          100.0%; Score 7; DB 20; Length 7; 100.0%; Fred R9 7 40.5; 5; 10dels
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                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB55781 standard; Peptide; 5 AA.
                                            Claim 1; Page 11; 16pp; English.
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99US-0134118.
99US-0150860
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Endothelial cell; haematopoietic cell; PDZ domain protein:
PL domain protein; leukcoyte activation; T cell surface receptor:
synapse formation; transmembrane neurotransmitter receptor:
autilmnuse disease; transplantation rejection, inflammation, allergy;
inflammatory bowel disease, ulcerative colitis, inflammation allergy;
astima; atopic dermatities atherosclerosis; cancer; infectious disease;
ischaemia; vasulitis; Crohn's disease.
Diplomical function of an ondothelia; cell or hematopoietic cell. The method involves introducing into a cell, an antagonist that inhibits binding between a PDY protein and a PL protein. The inhibits is used to troat a disease mediated by hematopoietic cells, e.g. autoimmuse disease. It may also be used to prevent transplantation rejection of a solil organ transplant. The method may also be used in the troatment of inflammation, allery, inflammatory bowel diseases, of existing colling indiations around diseases (e.g. theumatoid arthritis, multiple sclerosis, autoimmuse diseases (e.g. theumatoid arthritis, multiple sclerosis, autoimmuse disease) disease, isohemation rejection), atherosclerosis, reacers, infectious diseases, isohemia, vasulitis and Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New inhibitors of binding of a PDZ protein and PL protein for inhibiting T cellomediated response by hemotopoletic cells, or for treating diseases characterized by inflammatory and humoral immune
                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                        57.1%; Score 4: DB 22; Length 5: 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                     100.0%; Preu. ...
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The present invention relates to a method for modulating a biological function of an endothelial cell or haematopoletic cell, comprises introducing into a cell an antagonist that inhibits binding between a post domain protein and a PL domain protein to result in inhibition of leukocyte activation. The present sequence is a core sequence of a PL domain protein and a T cell surface receptor). Plu domains of proteins are named after three prototypical proteins: PSD4. Drosophila large disc protein and Zonia Gordudin introceins: PSD4. Drosophila large disc protein and Zonia Gordudin interactions: PSD4. Drosophila large disc protein and Zonia Gordudin interactions: PSD4. Drosophila large fluorities in symapse formation by organising transmembrane neurotransmitter receptors fromth intracellular interactions: The inhibitors identified by the present invention can be used to treat a disease mediated by haematopoietic cells, e.g. autoimmune disease. Inflammation, allergy cell dermitis, psoriasis, respiratory balled diseases, uscentive collitis, clicitis, psoriasis, respiratory valledge diseases, e.g. asthma, attended clickers, insulin-dependent diseases. Hashirvey thyrodisis, multiple schematikis, a their desease (e.g. thematicious disease. The used in prevent transplantation rejection of a collicity collicy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      57.1%; Score 4: DP 22: Length 5: 100 0%; Pred. No. 7.8e+n6; Pred. No. 7.8e+n6; Indels 1ye 0; Mismatches 0; Indels
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99US-0134117.
99US-0134118.
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99US-0170453.
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                                                                                                                                                                                                                                                                                                                                                                                  solid organ transplant.
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Matches 4, Conserv
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The present invention relates to a method for modulating a biological function of an endochelial cell or hasmatopoletic cell. Comprises introducing into a real an antagonist that inhibits binding between a piz domain protein and a PL domain protein to result in inhibition of leukwayte activation. The present sequence is a core sequence of a PL domain protein and a protein antagonic of a PL domain protein and a contact and proteins are named after three proteinsing protein. For Jomain proteins are included in symapse formation by organising transmembrane neurotransmitter. The present invention can be used to treat a disease mediated by the present invention can be used to treat a disease mediated by harmacopoletic cells, e.g. autoimmune disease, inflammation, allery, harmacopoletic cells, e.g. autoimmune diseases, ulcerative colifies. Celletts, psoriasis, reliammatory bowei diseases, ulcerative colifies dermatitis, autoimune diseases (e.g. asthma), atopic dermatitis, autoimune diseases (e.g. rheumatoid arthitis, multiple selection), isohaemia, vasullitis and Crohn's lisease. The celletts infection, isohaemia, vasullitis and Crohn's lisease. The
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                                               Modulating a biological function of a hematopoletic cell for treating
                                                                an allergic response, or diseases mediated by immune system cells, complises introducing into the cell a PDS-PE interaction enhancer or
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                                                                                                                                       Disclosure, Page 95; 143pp; English.
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Conservative 0.
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200008-0196527.
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es 4. Conservat
               WPI, 2001-061214,707.
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11-APR-2000;
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                                                                                                       inhibitor
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11-APR-2000;

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AAB58024
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Endothelial cell; haematopoietic cell; PDZ demain protein;
PL domain protein; leukocyte activation, T del surface receptor;
synapse formation; transmembrane neurotransmitter receptor;
autoimmune disease; transplantation rejection, inflammation; alleryy,
inflammatory bowel disease, ulcerative colitus; helitis; psoriasis;
asthma; atopic dermatitis; atherosclerosis; concer; infectious disease;
                                                                                                                                                                                                                                                                                                                                         Disological function of an endothelial cell of hematopoietic cell. The method involves introducing into a cell, as actasmist that inhibits binding between a PDS protein and a PL protein. The inhibitor is used to treat a disease mediated by hematopoietic cells, e.g. autoimmune disease. It may also be used to prevent transplantation rejection of a solid organ transplant. The method may also be used in the treatment of inflammation, albergy, inflammatory bowel itseases, ulcerative colitis, lieitis, psoriasis, asthma, atopic definitis, autoimmune diseases (e.g. theumatoid atthitus, multiple sclerosis, incerative diseases diseases, lasting transplantation thyroidiis, osteoarthritis, graft rejection, transplantation rejection, inherositatiss, dancers, infectious diseases, ischemia, vasulitis and rohm's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sdet in
                                                                                                                                                       Modulating a biclogical function of an endoth-dial rell or hematopoietic cell, useful for treating autoimume diseases and intectious diseases, by administering an anta-onist that inhibits binding between a PDZ protein and a PL protein.
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                                                                                                                                                                                                                                                                              Disclosure; Page 87-94; 141pp; English.
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                     (ARD-1-) ARBOR VITA CORP.
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14 - FEB-2800;
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function of an endothelial cell or haematopoletic cell, comprises introducing into a cell an antagonist that inhibits binding between a polyadmain protein and a PL domain protein to result in inhibition of pD2 domain protein and a PL domain protein to result in inhibition of leukooyte activation. The present sequence is a core sequence of a PL domain protein (a T cell surface receptor). PD2 domains of proteins are mamed after three protetypical proteins. PD2 domains of proteins are protein, and conclain proteins. PD2 domain proteins are included in synapse formation by organising transmembrane neurotransmitter receptors. Exhogin intracellular interactions. The inhibiture identified by the present invention can be used to treat a disease modified by the present invention can be used to treat a disease modified by the present cells, e.g. autoimmune disease, inflammation, allergy (e.g. drug allergies), inflammatory bowel diseases, uncertaive colitis, leftils, psoriasis, respiratory allergic diseases (e.g. asthma), alorgic dermattis, autoimmune disease. Hashmoto thyroidilis, contents in the dependent diabetes, Hashmoto thyroidilis, contents in the dispendent diabetes. Hashmoto thyroidilis,
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PL domain protein; leakecyte activation; T cell surface receptor;
synapse formation; transmembrane neurotransmitter receptor;
autoimmune disease; transplantation rejection; inflammation; allery;;
inflammatory bowel disease; ulcerative colitis; ileltis; psoriasis;
asthma: atopic dermatitis; atheresclerosis; cancer: infertious disease;
ischaemia; vasulitis; Crohn's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                            New inhibitors of binding of a PDZ protein and PL protein for inhibiting T coll-mediated response by homatopoietic cells, or for treating diseases characterized by inflammatory and humoral immune responses, e.g. inflammation, cancer
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Matches 4; Conservative 0
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11-APR-2000; 2000US-0196460.
11-APR-2000; 2000US-0196527.
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                                                                                                                                             (ARBO-) ARBOR VITA CORP.
                                                                                                                                                                                                                                                                                                                             WPI: 2001-025003/03.
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The present invention relates to a method for analyzating a biological function of an enduthelial cell or basmatopoietic cell, comprises introducing into a cell an antagonist that inhibite binding between a producing into a cell an antagonist that inhibition of the domain protein and a PL domain protein to result in inhibition of leuknoyte activation. The present sequence is a core sequence of a PL domain protein and a PL domain protein and atter three prototypical proteins. PSPS, Drosophila large disc protein and Zonnia vocludin i proteins. PSPS, Drosophila large disc protein and Zonnia vocludin i proteins. The inhibitors identified in synapse formation by organising transmembrane neurotransmitter receptors through intracollular interactions. The inhibitors identified by the present invention can be used to treat a disease mediated by hammatopoletic cells, e.g. autoimmune disease, inflammation, allergy (e.g. dermaticis, psorlasis, respiratory allergic diseases (e.g. asthma), atopic dermatitis, usulin-dependent diseases (e.g. rheumatod arthritis, multiple sclenosis, isulin-dependent diseases, inflammatory autoricus diseases (e.g. viral infection), ischaemia, vasulitis and Crohn's disease. The inhibitors can also be used to prevent transplantation rejection of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                              Modulating a biological function of a hematopoletic cell for treating an altergic response, or diseases mediated by immune system cells, corprises introducing into the rell a FPC-FF (interaction enhancer or inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hematopoletic cell: PDZ: PL: autoimmune disease; inflammation; allergy; asthma; multiple sclerosis; cancer; infection.
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100.0%; Pred. No.
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                99US-0134114.
99US-0134117.
99US-0134118.
99US-0160860.
                                                                                                                                   14-FE9-2000) 26000S-6182296.
11-APR-2000; 20900S-0195460
11-APR-2000; 20900R-0195527.
                                                                                                                     14-JAN-2000; 2000US-0176195.
                                                                                    99US-0162498
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Best Local Similarity 100.0
These 4: Conservative
                                                                                                                                                                                                           (ARBO-) ARBOR VITA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         solid organ transplant.
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21-OCT-1999;
                                                                                    50-0CT-1999;
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PL domain profeso, tenkonyte autivation; T cell surface receptor: synapse formation; transmembrane neurotransmitter receptor: autoimmune disease, transplantation rejection; inflammation, allergy; inflammatory bowel disease; ulcerative colitis; ileitis; psoriasis; asthma, atopic dermatitis; atherosclerosis; cancer; infectious disease; ischaemia; vasulitis; Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a solid organ fransplant. The method may also be used in the treatment of inflammation, allergy, inflammatory bowel diseases, ulcerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               collits, lefitis, psorfasis, asthma, atopic dermatifis, autoimmune diseases (e.g. rheumatoid arthritis, multiple solerosis, insulin-dependent diabetes, Hashimoto thyroiditis, osteoarthritis, graft rejection, afherosoletosis, can ers, infectious diseases, ischemia, vasulitis and Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a new method for modulating a biological function of an endothelial cell of hemotopoletic cell. The method involves introducing into a cell, an antagonist that inhibits binding between a FDZ protein and a FL protein. The inhibitor is used to treat a disease mediated by hematopoletic cells, e.g. autoimmune disease. It may also be used to prevent transplantation rejection of
                                                                                                                                                                                                                                                                                                                                                                                                   hematopojetic cell, useful for treating autoimmune diseases and
infectious diseases, by administering an antagonist that inhibits
binding between a FDZ protein and a FL protein
                                                                                                                                                                                                                                                                                                                                                                                Modulating a biological function of an endothelial cell or
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                                                                                                                                                                                     2000US-0176195.
2000US-0182296.
2000US-0196460.
                                                                                  99US-0134114.
99US-0134117.
99US-0134118.
                                                 12-MAY-2000; 2000W0-US13161
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99US-0162498.
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                                                                                                                                                                                                           14-FEB-2000;
11-APR-2000;
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                                                                                                    14-MAY-1999;
                                                                                                                       14-MAY-1999;
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29 - 001 - 1999;
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                23-NOV-2000
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Homo sapiens

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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ostecarhitis), atheroscierosis, cancers, intections diseases (e.g. viral infection). Ischaemis, vasullis and Chino's disease. The inhibitors can also be used to provent fransplantation rejection of a solid organ transplant.
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6. 7.8e++5;
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99US-0160860.
99US-0162498.
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                                                                                                                       99US-0134114
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2000mg-0144440
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              White Annual Page 1 - A2
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Matches
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The present invention relates to a method for modulating a biological function of an endothelial cell or haematopoietic cell, comprises introducing into a cell an antagonist that inhibits binding between a first domain protein and a Fi domain protein to result an inhibition of leukocyte activation. The present sequence is a core sequence of a Fi domain protein and a Fi domain protein sequence of a PZ domain protein and a First proteins. PSPS, prosphila large disc protein and Zennia protein by oldarising transmembrane neurotransmitter. In synapse formation by oldarising transmembrane neurotransmitter receptors through intracellular interactions. The inhibitors identified by the present invention can be used to treat a disease mailated by heamatopoietic cells, e.g. autoimmune disease, inflammation allery (e.g. drug alleques), inflammatory bowel diseases, inclammed alleris. Commattis, autoimmune diseases (e.g. estima) alterial contains a sclerosis, insulin dependent diabetes. Hashimoto thyroiditis.

Sclerosis, insulin dependent diabetes, Hashimoto thyroiditis.

Osteoritis, insulin dependent diabetes, Hashimoto thyroiditis.

Osteoritis, insulin dependent diabetes, intenticus disease. The conflator con also be used to prevent transplantation rejection is
Endothelial cell, haemalopoietic cell, FDZ domain protein;
FL Sozial, profeti, leukvoyte abliatica, T cell sufface receptor;
Synapse formation transmemblane neurotransmitter receptor;
synapse formation, transplantation rejection; inflammation; alleray;
autoimmune disease, transplantation rejection; inflammation; alleray;
inflammatory bowel disease; ulcerative colitis; iteltis; psoriasis;
astima, allej denatitis, albertasierosis, cancer; infectious disease;
ischaemia; vasulitis; groh's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Modulating a biological function of a hematopoietic cell for treating an allergic response, or diseases mediated by immune system cells, comprises introducing into the cell a PDZ-PL interaction enhancer or
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100.0%; Pic
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99US-0134117.
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14-7AN-2097; 200008-0176235.
14-888-2005; 200008-0186460.
11-APR-2009; 200008-0196457.
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                                                                                                                                                                      Homo sapiens.
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14-MAY-1999;
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3 KEYF 6
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AAB57616;

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The present invention relates to a new method for modulating a biological function of an endothelial cell of Pematopoietic cell. The method involves introducing income cell. The including between a PRZ protein and a PL protein. The including televation appropriate the including televation and a PL protein. The including is used to treat a disease mediated by hematopoietic cils, e.g. autoimmune disease. It may also be used to prevent transplantation rejection of a solid organ transplant. The method may also be used in the treatment of inflammation, allergy, inflammatory boxel issues, ulcerative colitis, ipsuidis, proteins dermatitis, autoimmune diseases (e.g. rheumatoid attritits, multiple ciletosis, insulin-dependent diabetes, Hashimoto thyroidits, csteoarth itis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    graft rejection, transplantation rejection, otherosclerosis, cancers, infectious diseases, ischemia, vasulitis and cobm's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Modulating a biological function of an endition of hell of hematopoletic cell, useful for treating autohomene diseases and infectious diseases. By administrating an antanomist that inhibits binding between a PDZ protein and a PL protein.
                                                                                                                                   Hematopoletic cell; PDZ; PL; autoimmune disease: inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57,1%; Soute 4; DR 22, Length 8; 100,0%; pred No 7 Herming tive 0; Mismatches 0; Indels
                                                                                                                                                   allergy, asthma, multiple sclerosis, cancer, infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatrhes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 87-94; 141pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB57616 standard; Peptide; 8 AA
            AAB55784 standard; Peptide; 8 AA
                                                                                                                                                                                                                                                                                                                  990S-0134117.
990S-0134118.
                                                                                                                                                                                                                                                                                                                                                                              9908-0170453.
                                                                                                                                                                                                                                                                                                                                                                                            zööcüs-öl/kly5.
Zööcüs-öl82296.
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200018-0146527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ARBO-) ARBOR VITA CORP.
                                                                                                     PDZ motif sequence #15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-080245,09.
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                                                                                                                                                                                                                                                                                                                                                                                           14-JAN-2000; 2
14-FEB-2000; 2
11-APR-2000; 3
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29-00T-1999;
13-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                          11-APR-2000;
                                                                                                                                                                                                                                                                       12-MAY-2000;
                                                                         07-MAR-2001
                                                                                                                                                                                                                                                                                                     14-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                   14 - MAY - 1999;
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                                                                                                                                                                                                                                          23-NOV-2000
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                                                                                                                                                                                Synthetic
                                            AAB55784;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lu PS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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AAB55784
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The present invention relates to a method for modulating a billogical function of an endothelial only of harmonicity cell computers introducing into a cell an antequentst Unat inhibits binding between a PDZ domain protein and a FL domain protein in inhibition of leukocyte activation. The present sequence is a core sequence of a PL domain protein (1 real) suiface receptors, PDZ domains of proteins are named after three proteingly proteins; PSDGS, Drosephila large discipancein and accordance of proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in syndry formation by organising transmentation in the contransmitter receptors through intravellular interactions. The inhibitors identified by the present invention can be used to treat a disease mediated by heamatopoietic cells, e.g. autoimmune disease, inflammation, alicryy (e.g. drug allergies), inflammatory bowel disease, ulcerative collitis, liellis, psochasis, respiratory allergic diseases (e.g. asthma), atopic dermatilis, autoimmune diseases (e.g. rheumatoid arthitis, autoimmune diseases (e.g. rheumatoid arthitis, autitiple solerosts, insulin-dependent diabetes, Hashimoto thyroiditis.
                                                                                                                                                               PL domain protein; leukocyte activation; T cell surface receptor; synapse formation; transmembrane neurotransmitter receptor; autoimmune disease; transplantation rejection; inflammation; allerny; inflammatory bowel disease; alcerative colitis; ileitis, psoriasis; asthma; atopic dermatitis; atherosclerosis; cancer; infectious disease; ischaemia; vasulitis; Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       osteparthritis) arbarocolerosis, cancers, infections diseases (e.g. viral infection), ischaemia, vasulitis and Crohn's disease. The inhibitors can also be used to prevent transplantation rejection of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sde:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New inhibitions of binding of a PDZ protein and PL protein for inhibiting I cell-mediated response by hematopoletic cells or for treating diseases characterized by inflammatory and humoral immune
                                                                                                                                 Endothelial cell; haematcpoietic cell, PDZ domain protein:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inde s
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                                                              31ycophorin C C-terminal core sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         responses, e.g. inflanmation, cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure: Page 93: 139pp: English.
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200008-0182296.
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990S-0134117.
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(first entry)
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Best Local Similarity 100.
Matches 4: Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ARBO-) ARBOR VITA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         solid organ transplant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WELL - 2002-125055 (03)
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                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-APR-2000;
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14-FFH-2000;
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The present invention relates to a method for modulating a biological function of an endothelial cell or hapmatop, orto cell compresses introducing into a cell an antagonist that inhibits binding between a PDZ domain protein and a PL domain protein. I result in inhibition of leukocyte activation. The present sequence is a core sequence of a PL domain protein (a T cell surface receptor). BZ domains of proteins are named after three protecypinal proteins: PSL-45, Prosophila large disc protein and Zounda Occludin 1 protein. PDZ domains of proteins are involved in symmetry formation by organising transments one neurotransmitter receptors through intraceflucial interactions. The inhibitors identified by the present invention can be used to treat a disease mediated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         håematopotetic cells, e.g. autoimmune disease, inflammation, allerjy, (e.g. duug allergies), inflammatory bowel diseases, ulcerative collitis, ilettis, psoriasis, respiratory allergic diseases (e.g. asthma), atopic dermatitis, autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, insulin-dependent diabetes, Hashimoto thyroiditis, osteoarthritis), atherosclerosis, cancers, infectious diseases (e.g. viral infection, ischaemia, vasulitis and cohn's disease. The
                                                                                                                                                                                                                                                                                                     Endothelial cell, haematopoietic cell, FDZ domain protein, PDZ domain protein, leuchtoric techtoric surface ieceptor, synapse formation; transmembrane neurotransmitter receptor; autoimmune disease, transplantation rejection, inflammation; allergy; inflammatory bowel disease, ulcerative colitis; inflammation powel disease, ulcerative colitis; ilettis; psoriasis; asthma; atopic dermatitis; atherosclerosis; canver; infectious disease; ischemnia; vasulitis; Crohn's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modulating a biological function of a hemator-letic cell for treating an allergic response, or diseases mediated by immune system cells, comprises introducing into the cell a PDZ-PL interaction enhancer or
                                                                                                                                                                                                                                                                 Glycophorin C C-terminal core sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure: Page 96: 143pp; English.
                                                                                                                                           AAB58026 standard: Pertide: 9 AA
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200008-0196527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MPI; 2001-061214,007
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29-00T-1999;
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2 KEYF 5
                                      4 KEYF 7
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                                                                                                                                                                                    AAB58026;
                                                                                                  RESULT 13
                                                                                                                          AAB58026
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AARS9104-264 are immunogenic 9mer paptides that contain a BLA-ALLI binding modif. These peptides bind HDA-AZ I and have a binding attituity of at least 1% as compared to a reference peptide (AAR71278), AARS922S has an incompared to a reference peptide (AAR71278), AARS922S protection. The peptides of the invention can induce systems in Immibosytes which can react with target cells. They can be used for the treatment or prophyaxis of cancer, eg. prostate cancer or lymphoma, etc.
inhibitors can also be used to prevent transplantation rejection of a solid organ transplant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sake.
                                                                                                                                                                                                                                                                                                                                                               antigen; epitope; immunogenic target protein; PSA; HBVC; HBVS; EBV; HTVI; corr antigen; surface antigen; pharmaceutical composition; in vivo; ax vivo; therapeutic; diagnostic; MHC class I molecule; major histocompatability complex; HLA-A2:1; 9mer; 10mer; archer:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunogenic peptide(s) having an HLA-A2.1 binding motif - used for treatment or prophylaxis of cancer, virus infection or autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             57.18; Score 4; DB 15; Length 9: 100.08; Pred. No. 7.8e+05; rive 0: Mismatches 0: Indels
                                                                                                    0; Indels
                                                                            Length 8;
                                                                            57,1%; Score 4: DB 22; Le
100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                      Peptide fragment (1.0268) of P53 binds HLA-A2.1.
                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; Page 105; 138pp; English.
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                                                                             57,15,
100.0%; Pr
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9308-0073205.
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9308-0159184
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                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                               human leukocyte antigen.
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                                                                                            Local Similarity
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                                               8 AA:
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29 - NoV - 1993.
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                                                                                                                                       2 KEYF 5
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                                                                                                                                                                     4 KEYF 7
                                                 Sednence
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                                                                             Query Match
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Matches
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RESULT 15

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p53; Her-2; Neu, aa; amino acid: CTL: cytctesin T lymphocyte; target;
malignant cell: antigenic; varcine; immunisation: activation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ARB97517-P97544 are antifenic p53-derived peptides synthasised to be used in an assay for identifying peptides cap-ble of activating cytotoxic I nymbiocytes (CTLS) which specifically target malignant cells. CTL-activating peptides can be used in a vaccine for protecting against tumour cell formation. CTL-activated by the peptides will lyse tumour cells displaying specific peptides. Antibodies against CTL-activating peptides are useful for the identification of other similar compounds which may be useful for treating carcer or virally-infected cells, or for diagnosis The peptide and vaccines produced provide immunity to a high percentage of different ethnic groups, i.e. those with different file alleles.
                                                                                                                                                                                                                                                                                                                                                                                                                                              In vivo activation of tumbur-specific sytutuals T lymphasytes - by contacting with polypeptide(s) derived from human p53 or Her-2/Neu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       57.1%; Score 4; DB 17; Length 9; 100.0%; Pred, No. 7.8e+95; tive 0; Mismatches 0; Indels
                                                                                                          Antigenic peptide, corresp. to p53 aa 322-330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 94; 158pp; English.
        AAR97542 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                     94US-035558.
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Rest Local Similarity 100...
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                                        AAR97542;
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AAR97542
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